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OP 62 PERIODEDIS 612  Db 61 PERIODEDIS 71  RESULT  1 OSVYEZ PRELIMINARY. PRT; 84 AA.  AC OSWYEZ PRELIMINARY. PRT; 84 AA.  AC OSWYEZ PRELIMINARY. PRT; 84 AB.  AC OSWYEZ PRELIMINARY. PRT; 84 AB.  AC OSWYEZ PROSTATE SPECIFIC MEMBRANE ANTIGEN PSM (PRAMENT).  DF 61 -007 2000 (Tremble: 1.5). Carated  DF 61 -007 2000 (Tremble: 1.5). Last annotation update).  DF 61 -007 2000 (Tremble: 1.5). Last annotation update).  DF 61 -007 2000 (Tremble: 1.5). Last annotation update).  DF 61 -007 2000 (Tremble: 1.5). Last annotation update).  DF 61 -007 2000 (Tremble: 1.5). Last annotation update).  CC Mammalia; Estination (Control & S.)  RF 62 -007 2000 (Tremble: 1.5). Last annotation update).  DE 61 -007 2000 (Tremble: 1.5). Last annotation update).  DF 61 -007 2000 (Tremble: 1.5). Last annotation update).	NCBL_TaxID=9608; [1] [1] [2] [2] [2] [2] [2] [3] [2] [2] [3] [2] [2] [3] [2] [3] [3] [3] [4] [4] [4] [4] [4] [4] [4] [4] [4] [4	Mon Apr 9 11:14:16 2001 ue-08-403-803-2_copy_45_750.olig.rept
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77 01-ANN-198 (Trembirel 05 Cartaced)

77 01-ANN-198 (Trembirel 15 Last annotation update)

78 01-CANN-198 (Trembirel 15 Last annotation update)

79 01-CANN-198 (Trembirel 15 Last annotation update)

80 POLAT OR HOPSM.

81 POLAT OR HOPSM.

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Matches 50; Conservative 0;
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Best Local Similarity 100.0%; F
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BML, U7997] AMC51431 I.
BML, RP000254; AMC40067 1.
BML, RP000275; AM66759 1.
HEROSE, M28 0.01
INTERPRO, LEPRO01177;
PPAM, PPO02255 PA, 1.
Carbonypertidase; Mydrolase,
BROWENCE 752 AM: 04539 MM; 500915A3B9C71E41
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EMEL ARF050502; AAC39269.1;
HERDFS: M28.0103137;
HTTERPO: 15F0003137;
CARDOTPEPTIGMS: F07225: F0A.17
CARDOTPEPTIGMS: F07235 F0A.27
SEQUENCE: 731 AA; 84523 MF; AF77835;
Query Match 3.4%; Score 24; DB 4; Le Best Local Similarity 100.0%; Pred. No. 3s-16; Hatches 24; Conservative 0; Mismatches 0;
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TISSUE-PERSON
HEDLINE-993120AP PROOSE:
HEDLINE-993120AP Duth1-Carter R., Villanueva J.A., Gardner J.M.,
Haisted C.H., Ling E., Luth1-Carter R., Villanueva J.A., Gardner J.M.,
COYLE, J.T.;
COYLE, J.T.;
                                                                                                                                                                             SECURENCE FROM N.A.
TISSEP-LUNG CARTHOMA:
TISSEP-LUNG CARCITROMA:
TISSEP-LUNG CARCITROMA:
REDLINE-99185063; PubMed-10085079;
Repagalos M.N. Peefs J.M., Somers M., Fraiponts E., Ashton D.,
Gordon R.G.;
*Igolation and expression of novel human glutamate carboxypeptidases
with N-acetylated alpha-linked acidic dipeptidase and dipeptidyl
peptidase T westlytty.
peptidase T westlytty.
J. Biol. Chem. 274-8870-8483(1999).
MEMOLE MODIATO. CARSSEST.1:
**SEQUENCE 100 AA; 83591 NM; 040624D591ECF879 CRC64;
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Necular characterization and relation to glutamate carboxypeptidase
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NCE 752 AA; 84635 MW;
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Sciurognathi; Muridae; Murinae;
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OLSJÖS PRELLIMINARY; PRT; 29 AA.

OLSJÖS PRELLIMINARY; PRT; 29 AA.

OLSJÖS PRELLIMINARY; PRT; 29 AA.

OLSJÖS PROST, PRT; 29 AA.

DT 01-NOV-1996 (TEMBLIFA! 01, Last sequence update)

DT 01-NOV-1996 (TEMBLIFA! 15; Last annotation update)

DT 01-NOV-1996 (TEMBLIFA! 15; Last sequence update)

DT 01-NOV-1996 (TEMBLIFA! 15; Last sequence update)

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HOURS PROSTATION (TEMBLIFA! NATION (TEMBLIFA! 15)

RE NOUTH PROSTATION (TEMBLIFA! TEMBLIFA! 15)

RE SEQUENCE FROM N. A.

RC TISSIDE-PROSTATIS

RX MEDLING-95188188; PubMed-7882149;

RX MEDLING-95188188; PubMed-7882149;

RX MEDLING-95188188; PubMed-7882149;

RX TISSIDE-PROSTATIS

RX MEDLING-95188188; PUBMEd-7882149;

RX TISSIDE-PROSTATIS

RX TISTIO (TEMBLIFA! 15)

RX TISTIO (TEMBLIFA! 15)

RX ALTERIATIVELY SPLICED VALIDATE OF POSITION W. D.

RX TISTIO (TEMBLIFA! 15)

RX MEDLING-95188188; PUBMEd-7882149;

RX TISTIO (TEMBLIFA! 15)

RX ALTERIATIVELY SPLICED VALIDATE OF POSITION W. D.

RX ALTERIATIVELY SPLICED VALIDATE OF POSITION W.
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P70627:
P7062
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TISSUE-PROSTATE:
MEDLINE-95188180: pubMed-7882349;
MEDLINE-951881818: pubMed-7882349;
MS.L., Nuang I.P., Fair W.R., Fowell C.T., Heston W.D.;
Su.S.L., Standard of prostate specific membrane antigen
Alternatively spiliced variants of prostate specific membrane antigen
RNA: ratio of expression as a potential measurement of progression. ';
Cancer Res. 55:1411-144(1995).
EMBL; S76578. AAB3750.2; '.
SEQUENCE 29 AA: 3455 MW; IDCSOEAF638670F3 CRC64;
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Carter N. E., Feldman A. R., COYLe J. T.;
Probletic specific membrane antiquo is a hydrolase with
Probletic specific membrane antiquo is a hydrolase with
pharmacologic characteristics of a outopeptidase.";
Proc. Matl. Acad. Sci. U.S.A. 33.749-753(1996).
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TISSUE-BBAIN;
REDLINE-88:66524; PubMed-9501243;
Luthi-Carter R., Berger U.V., Barcask A.K., Enna M., Coyle J.T.;
Isolation and expression of a rat brain cDNN encoding glutamate carboxypeptidase TU.Y.
Action and expression of a Price Carboxypeptidase TU.Y.
Proc. Matl. Acad. Sci. U.S.A. 95:3215-3220(1998).
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STRAIN-SPRAQUE DAWLEY: TISSUE-BRAIN:
MEDLINE-98041505; PubMed-9375657;
BEddega T., Turi T., Wroblevska B., She D.,
Weale J.H.
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TISSUE=BRAIN:
Luthi-Carter R., Berger U.V., Barozak A.K.,
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from a rat hippocampal cDNA library ;
J. meurochem. 68:2276-2277(1997).
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pred. No. 5.5e+29;
0; Mismatches 0;
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Catarrhini; Hominidae; Homo.
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                             Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                  PRT;
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320 GAVEPDRYVI 329 ||||||||| 360 GAVEPDRYVI 369

O53737 PRELIMINARY: PRT: 485 AA.
053737; PRELIMINARY: PRT: 485 AA.
053737; O1.NOV-1996 (TYEMSLEVEL. O1. Created)
01.NOV-1996 (TYEMSLEVEL. O1. Last sequence update)
01.OCT-2000 (TYEMSHLEVEL. O1. Last sequence update)
01.OCT-2000 (TYEMSHLEVEL.) IS. Last sequence update)

Streptomyces lipmanii (Streptomyces alboniger).
Booteria: Firmioutes; Actinobacteria; Actinobacteridae;
Actinomycetales; Streptomycineae; Streptomycetaceae; Streptomyces.
NBP\_TWANID-1892;

Query Match
Best Local Similarity 100.0%; F
Matches 10; Conservative 0;

Score 10; DB 11; Pred. No. 0.23; 0; Mismatches 0

0; Indels 0;

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Length 745;

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Mammalia; Eutheria: Primates; Catarrhini; Hominidae: Homo NCDI\_TaxID=9606;

signment prom M., Neefs J.M., Somers M., Fraiponts E., Ashton D.,
C TISSUB-MSALL INTESTINE:
PANGALOS M.N., Neefs J.M., Somers M., Fraiponts E., Ashton D.,
A Gordon R.G.,
Tisolation and expression of novel human glutamate carboxypeptidaes
of with M-acetylated alpha-linked acidic dipeptidase and dipeptidyl
peptidase IV actvity.
peptidase IV actvity.
EMBL. AND 1271; CAB3956.1;
EMBL. AND 1271; CAB395

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STRAILM-SPRAGEE DAWLEY: THE STRAILM-SPRAGE N.C., Rinaldo P.,

Devarsjan P., Sun A.G., Dawson P.A., Ananthanarayanan M.;

J. Biol. Chem. 0:0-0(1997).

EMBL. AP00921; JAB87644.1:

INTERPRO: 12800117:

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EMBL: AL1386708 COART-592 J. F.
ESPOEMEN TO 3 AA: 76925 MM; 28814DFF11F4B204 CRC64;

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BE TO SH: PJ3817.

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CH AGENA VIUSES, NO RNA STAGE: POXVIIIdae; Entomopoxviriae;

CO VIUSES: AGENA VIUSES, NO RNA STAGE: POXVIIIdae; Entomopoxviriae;

CO NCOLIMINITES 3.101.

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Arabidopsis thaliana (House-ear cress).
Eukaryota: Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;
Magnollophyta; eudicoryledons; core eudicots; Rosidae; eurosids II;
Brassicales; Brassicoceae; Arabidopsis.
KRPL\_TaxID=3702;

SEQUENCE PROM N.A.

Obermaker B., Ottenwaelder B., Duchemin D., Zeitler K., Meves H.W.,
Lemcke K., Mayer K.F.X., Quetter F., Salanoubat M.;
Lemcke K., Mayer K., Mayer K., Mayer K., Mayer K., Mayer M.;
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15016CCF70E5EBB0 CRC64

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STRAIN-TYCSON;
STRAIN-TYCSON;
MEDILINE-99.02612; PubMed-9847159;
Alcond C.L., Thlann E.R., Lu Z., Ona E., Rutish G.F., Rock
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SEQUENCE FROM N.A.

Afonso C.L., Tulman E.R., Lu Z., Oma E., Kutish G.P., Rock D.L.:
Submitted (RMT-1998) to the EMBL/GenBank/DBBJ databases.
EMBL, AP663865, AC97675.1;
SEQUENCE 1306 AA; 153649 MW; PC3868A8292C5904 CRC64;

Query Match 1.3%; Best Local Similarity 100.0%; Matches 9; Conservative 82 NYISIINED 90 |||||||| 160 NYISIINED 168 Score 9; DB 12; Length 1306; Pred. No. 4.5; 0; Mismatches 0; Indels

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Neisseria meningitidis (serogroup B) Bacteria: Proteobacteria: beta Subdivision: Neisseriaceae; MCDI\_TaxID+91:

Neisseria.

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NA MCDORLI I., HUTCHEN P. PROTECT J.D., MEADON W.C. Exchima K.A.,

NA MCDORLI I., HUTCHENCY R. PROTECT K.D., GATFER M.,

NA STEWART A.M., CORTON M.D., PPART M.S., PHILLIPS C.A., RICHARDSON D.,

NA Heidelberg J., Sutton G.G., Fleichmann R.D., Eisen J.A., White O.,

NA Salbberg G.L., Smith N.O., Venter J.C., Francer C.M.,

R. *Evidence for lateral gene transfer between Archaea and Bacteria from

RT genome sequence of Thermotoga maritima.';

NA STATEMENT OF THE STATEMENT OF THE PROTECT CONTROL OF THE PROT
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Best Local Similarity 100.0%;
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Best Local Similarity 100.0%; F
Matches 8; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Science 287:1809-1815(2000).
EMBL, AE003518; AAP42029:1; *.
TIGR, NMB1681; *.
Rypothetical protein.
SEQUENCE 141 AA; 15499 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 396 RLLQERGV 403
||||||||
40 RLLQERGV 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         031229 PRELIMINARY: PRT: 260 AA.
031279.998 (TREMBLEST. 05, Created)
01-7AN-1998 (TREMBLEST. 05, Least sequence update)
01-007-2000 (TREMBLEST. 15, Last senotation update)
MOLYBRATE SIBLOIMS PREIDLASSIC PROFISI.
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Kim S.O., Hyun C.G., Suh J.W.;
Fefficiant cloning of the genes for RNA polymerase sigma-like factors from Actinomycetts.;
J. Microbiol. Biotechnol. 8:280-283(1998).
EMBIL, A007193 JAC2665.1:
-UNITERE 0.590-00431;
PRIMITER 10.1

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||||||||
122 QAAAETLS 129
SEQUENCE FROM N.A. (1910) G., Nick P., Brandsch R., Schubach B., Mennidez C., Otto A., 1910) G., Nick P., Brandsch R., Schubach B., Mennidez C., Otto A., 1910) G., Nick P., Brandsch R., Schubach B., Mennidez C., Otto A., 1910) G., Nick P., Brandsch R., Schubach R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Arthrobacter nicotinovorans.
Plasmid pAO1.
Bacteria; Firmicutes; Actino
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|||||||
150 RLQDFDKS 157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bacteria; Firmicutes; Actinobacteria; Actinobacteridae;
Actinomycetales; Micrococcineae; Micrococcaceae; Arthrobacter.
KPL_TexID=29320;
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i; Pred. No. 6.1;
0; Mismatches
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; Pred. No. 11;
0; Mismatches
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o. 11;
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Best Local Similarity 100.0%;
Matches B; Conservative
       SEQUENCE FROM N.A.
STRAIN-CV. COLUMBIA;
Waterston R.;
Submitted (MAY-1999) (
[5]
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53 FTEIASKF 60
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||||||||
29 LLEKMGGS 36
                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
STRAIN-CV. COLUMBIA;
Waterston R.;
Submitted (APR-1999)
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09XIA9 PRELIMINARY: PRT: 280 AA.
09XIA9 (TERBLICAL 12, Created)
01-NOV-1999 (TERBLICAL 13, Last sequence update)
01-NOV-1999 (TERBLICAL 13, Last annotation update)
F10A2 10 FOR FILE (UPMATUE ENTIMARY)
F10A2 10 FOR FILE (UPMATUE ENTIMARY)
F10A2 10 OR AT4GO7570
Arabidopsis challana (Mouse-ear cress)
F10A2 10 OR AT4GO7570
Arabidopsis challana (Mouse-ear cress)
F10A2 10 OR AT4GO7570
Arabidopsis challana (Mouse-ear Cress)
F10A2 10 OR AT4GO7570
F10A2 10
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Plasmid.
SEQUENCE 260 AA;
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STRAIN-GV. COLUMBIA;
JOSHU C., Bauer C., Hotic M.;
The sequence of A. thaliana
Submitted (MAY-1999) to the E
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Submitted (MAY-1999) to
[2]
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STRAIN-CV. COLUMBIA;
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(MAY-1999) to
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0B28703B77E7981B CRC64;

us-08-403-803-2\_copy\_45\_750.olig.rspt

Page 36

Score 8; DB 2; pred. No. 11; 0; Mismatches

Length 260;

0; Indels

0; Gaps

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the

EMBL/GenBank/DDBJ databases

Sequencing Project."; the EMBL/GenBank/DDBJ

databases

F10A2. ; EMBL/GenBank/DDBJ

databases

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RESULT 17

OSM138

ID OSM138

AC OSM138:
DT 01-NOV-1999 (TEDRELTE1. 12, Created)
DT 01-NOV-1999 (TEDRELTE1. 12, Last sequence update)
DT 01-NOV-1999 (TEDRELTE1. 12, Last sequence update)
DT 01-NAY-2000 (TEDRELTE1. 13, Last sequence update)
DE SEPERITORIEVPUTRESCINE SOC TRANSPORTER, PERMEASE PROTEIN.
GN TMAIJORG MATICINA.
OC BACTERIA TREINDOGGALES; THETMOCOGA.
OX NCD1.TAXID-3356.
DE 11 UDDAGE FROM N. A.
SEMAINAESS / DSM 1109.
RX MEDLINE-99287316; PubMed=10360571;
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1.1%; Score 8; DB 4;
Best Local Similarity 100.0%; Pred. No. 7.1;
Matches 8; Conservative 0; Mismatches
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OUNTS:
OUNTS:
OUNTS:
OUNTS:
OL MAY-2000 (TEERBLEE), 13. Last sanotation update)
OL MAY-2000 (TEERBLEE), 13. Last sanotation update)
HYPOTHETICH, 17.1 KDA PROTEIN (FRAGHENT).
HYPOTHETICH, 18.1 HUMBON, 18.
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TISURS-UTERUS, Obermaier B., Mewes H.W., Gassenhuber J., Wiemann S.;
Submitted (AUG-1999) to the EMBL/GenBank/DDBJ databases.
EMBL, All0260 CABS3701.1; -.
BMBL, All0260 CABS3701.1; -.
NY TEB
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165 AA; 17070 MW;
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RESULT 23

PSI-06

PSI-07

PSI-07

PSI-08

AC PSI-06

PT 01-MAY-1997 (TERBLETA) .03. Creaked)

DT 01-MAY-1997 (TERBLETA) .03. Last sequence update)

DT 01-MAY-1997 (TERBLETA) .15. Last annotation update)

DE 1911-181-191 (TERBLETA) .15. Last annotation update)

DE SIHILAR TO HUMAN PROSTRATE-SPECIFOR MEDIANE ANTIGEN.

GN R57.1.

OC ENDARYOLA; Metazos; Nematoda; Chromadora; Rhabditida; Rhabditoides;

OC ENLARYOLA; Metazos; Nematoda; Caenorhabditis.

NEMB_TaxIDe6339;

RN SEQUENCE PROM N.A.

RC STALIMENISTOL N2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 22

OL382

ID 01382

PRELIMINARY, PRT; 694 AA.

AC 01382

DT 01-JAN-1998 (TTERBLEL) 05, Created)

DT 01-JAN-1998 (TTERBLEL) 05, Last sequence update)

DT 01-JAN-1998 (TTERBLEL) 13, Last annucation update)

DE MOLTBODOTERN LOPACTOR BIOSYNTHETIC PROTEIN

CRIADOTERN COPACTOR BIOSYNTHETIC PROTEIN

RA DORED D H, Kinghorn J R.;

RA DORED D
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Best Loc
Matches
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Best Local Similarity 100.0%; F
Matches 8; Conservative 0;
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Best Local Similarity 100.0%;
Matches 8; Conservative 0;
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GDALGE S.F., SMITH J. KARAH G.J.M.M., Millar L.J., Heck I.S.,

GDALGE S.F., SMITH J. KARAH J.

BAOL Chem O.G. (197)

J. BAOL Chem O.G. (197)

INTERPRO; IPPRO0235)

INTERPRO; IPPRO0235)

FPAN; PPO1467; MOAL, MITB_PQGE; 1.

FRAN; PPO1567; MOAL, MITB_PQGE; 1.

FROSITE: PS01305; MOAL, MITB_PQGE; 1.

FROSITE: PS01305; MOAL MITB_PQGE; 1.
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STRAIN=CV. COLUMBIA;
Waterston R.;
Submitted (JUN-1999) to t
[6]
                                                                                                                                                                                                                                                                                                                                                                                                                 260 KILEKMGG 267
||||||||
673 KILEKMGG 680
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|||||||
176 EPDRYVIL 183
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Bacterla: Proteobacteria: gamma subdivision; Alteromonadaceae;
Alteromonas.
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SEQUENCE FROM N.A.

SEQUENCE PROM N.A.

SUBMILITED (MR.7200) to the EMBL/GenBank/DDBJ database Submitted (MR.7200) and the EMBL/SEQUENCE (ASB1137.1).

EMBL/ALIGISOS (ASB1137.1).

SEQUENCE 280 AA; 31115 MH; AB7FECF4E1870196 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 8; DB 3;
Pred. No. 28;
O; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 8; DB 2; Pred. No. 13; 0; Mismatches
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Pred. No. 12;
0; Mismatches
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          RESULT
Q9M1E4
ID Q9
AC Q9
DT 01
DT 01
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Best Local Similarity 100.0%; I
Matches 8; Conservative 0;
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Best Local Similarity 100.0%; Matches 8; Conservative 0;
09MLEA, PRELIMINARY; PRT; 989 AA.
09MLEA;
01-CCT-2000 (TrEMBLrel. 15, Created)
01-CCT-2000 (TrEMBLrel. 15, Last sequence update)
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413 WDAEEFGL 420
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STRAIN-BRISTOL N2:

OZERSKY P., Schaller M.;

*The sequence of C. elegans of

Submitted (MAR-1999) to the E
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STRAIN-BRISTOL N2;
Waterston R.;
Submitted (FEB-1997) to t
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STRAIN=BRISTOL N2;
Favello T., Rifkin L., C
Submitted (PEB-1997) to
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Submitted (MAR-2000) to the EMEMBL; AC00656; AAF39882.1; -.
SEQUENCE 549 AA; 65212 MW;
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STRAIN-BRISTOL N2;
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SEDUENCE FROM N. A.
STRAIN-BEIGEL N2.
Whaterston B.D. 12.
Whaterston B.D. 19.
SUBmitted (ABR-1997) to the EMBL/GenBank/DDBJ dat
EMBL: U8819: AAB52660.1;
HEROPS: MAD 1010: -
HEROPS: MAD 101: -
INTERFRO, IPRO01317;
PPAMS: PRO7225; PA. 1.
SEQUENCE 751 AA: 83563 MH; 1A08A44974PA29C7 (
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OSSEM:
OSS
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Elbazyota, Metason, Memacoda; Chromadoren; Rhabditida; Rhabditoidea
Rhabditidae; Metason, Memacoda; Caenorhabditis,
NGL_TaxID=6239;
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STRAIN=BRISTOL N2;
MEDLINE=99069613; PubMed=9851916;
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o the EMBL/GenBank/DDBJ
Score 8; DB 5; Pred. No. 31; 0; Mismatches
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Pred. No. 23;
O; Mismatches
                                                                                                                                                                                                                                                1A08A44974FA29C7 CRC64;
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EMBL; AB025228; BAA95201.1; -. Transferase. SEQUENCE 1575 AA; 176792 MW;

772A26E4D7C2E543 CRC64;

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RY STREAM-CHALLS; pubMed=153435;
RX HELLIN-2276537; pubMed=153435;
RX HELLIN-2276537; pubMed=153435;
RY Jentification of a gene, reg, which regulates expression of RT Jentification of a gene, reg, which regulates expression of RT Joseph Library 134, 1357-3586(1992).

DR HELLIN-2343; MC448431;
DR HELLIN-2343; MC448431;
DR HELLIN-23458531;

RESULT 28
Q9RWZ7
LD Q9RWZ7
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Best Local Similarity 100.0%; F
Matches 8; Conservative 0;
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OSCH3;

O1-CCT-2000 (Tremstrel. 15, C
O1-CCT-2000 (Tremstrel. 15, L
O1-CCT-2000 (Tremstrel. 15, L
OLCCT-2000 (Tremstrel. 15, L
OLCCT-2000 (Tremstrel. 15, L
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Best Local Similarity 100.0

Matches 8; Conservative
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054178 (054247;
01-NOV-1996 (TERMELTEL 0)
01-NOV-1996 (TERMELTEL 0)
01-NOV-1996 (TERMELTEL 0)
01-NAY-2000 (TERMELTEL 1)
GLUCOSYLTRANSFERASE.
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HYPORETICAL 108.4 KDA PROTEIN.
PSP21.1210.
Arabidopais thaliana (House-ear Cress).
Eukaryota: Viridiplante; Embrophyta; Tracheophyta; Spermatophyta;
Magnollophyta; eudicoryledons; core eudicots; Rosidae; eurosids II;
Brassladies; Brassladocae; Arabidopsis.
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||||||||
545 RRGIAEAV 552
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JOrdan N., Bangert S., Wiedelmann R., Voss H., Unseld M., Heves H.W.,
Lemcke K., Mayer K.F.X., Quetier F., Salanoubat M.;
submitted (FEB-2000) to the EMBL/GenBank/DDBJ databases.
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SERVINNEE FROM N.A.
SERVINNEE STRAIN-CHARLES
MEDILINE-96157064; pubMed-8586195;
VLOKERMAN N.H., Sulavik N.C., Clevell D.B.;
VLOKERMAN N.H., Sulavik N.C., Clevell D.B.;
Photecular analysis of Streptococcus gordonii glucosyltransferase
phase variants. p
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STRAIN-MYCCLOST;
STRAIN-MYCCLOST;
Pujisara T., Hoshino T., Ooshima T., Sobue S., Hamada S.;
Purification, characterisation, and molecular analysis of the encoding glucosyltransferses From Streptococcus oralis.;
Infect. Immun. 68: 445-2481/2009).
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EU ARADIAOpaia sequencing project;

EU ARADIAOpaia sequencing project;

EUBL: AL13667EB-2000) to the EUBL/GenBank/DDBJ databases.

EUBL: AL13667EB-2000;

EUBL: AL1
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Bacteria; Firmicutes; Bacillus/Clostridium group; Streptococcaes.
Streptococcus.
Streptococcus.
INI_TaxID=1302;
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roteria: Pirmicutes: Bacillus/Clostridium group: Streptococcus.
reptococcus.
pl_TaxID=1303;
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Pred. No. 40;
0; Mismatches
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Pred. No. |
O: Mismatch
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Best Local Similarity 100.0%; Pred. No. 62;
Matches 8; Conservative 0; Mismatches
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P SCHEPHOCOCCUS SALLVARIOUS N.C. 25975 Possesses at least two gen of the control o
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Best Local Similarity 100.0%; 1
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999873 PRELIMINARY:
099873:
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01-MAY-2000 (TEMBLICEL 1
01-MAY-2000 (TEMBLICEL 1
CG14505 PROTEIN.
CG14505 PROTEIN.
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1059 AFLDELKA 1066
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Bacteria:
Streptococcus:
Streptococcus:
NCPI_TaxID=1304:
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1091 AFLDELKA 1098
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11 APGVKSY 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'Genome Sequence of the Radioresistant Bacterium Deinococcus radiodurans RI. 577(1999).
Science 286.1571-1577(1999).
BMBL. ABROUSID. AAP10098.1: -
TOR. ABROUSID. AAP10098.1: -
TOR. ABROUSID. AP1097 NP; DAIBBOOBA42E439A CRC64:
SEQUENCE 66 AA. 8797 NP; DAIBBOOBA42E439A CRC64:
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STRAIMERS, STRAIMERS, A.

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MEDIL
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01-MAY-2000 (TREMBLAR). 13. Last sequence update)
01-UN-2000 (TREMBLAR). 14. Last annotation update)
HRD7HETICAL 8.8 KDA PROTEIN.
DR0518.
Drosophila melanogaster (Fruit fly).
Bukaryota: Metazoa: Arthropoda: Trachesta: Hexapoda: Insecta:
Preryota: Neptera: Endopteryota: Diptera: Brachycera: Muscomorpha
Ephydroidea: Drosophilidae: Drosophila.
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Bacteria; Thermus/Peinococcus group; Deinococcales; Deinococcus
INCBL_TaxID=1299;
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                                                                                                                                                                                          Created)
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Pred. No. 62;
O; Mismatches
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RE SEQUENCE FROM N. R.A.

RE SEQUENCE FROM N. R.A.

RE SETTION C.G., Scherrer S.E., Michael M.D., Zhang G.A., Gocompton S.N.

RA Mannathides P.G., Scherrer S.E., Michael M.D., Zhang G.C., Chen L.X.

RA SETTION C.G., Wortman J.R., Yandell M.D., Zhang G.C., Chen L.X.

RA SETANDON C.G., Wortman J.R., Yandell M.D., Zhang G.C., Chen L.X.

RA SETANDON C.G., Wortman J.R., Yandell M.D., Zhang G.C., Pallown C.R., Miklos G.L.G.,

RA MANN K.H., Doyle C., Baxeer E.G., Hell G., Welson C.R., Miklos G.L.G.,

RA MANN K.H., Doyle C., Baxeer E.G., Hell G., Welson C.R., Miklos G.L.G.,

RA MANN K.H., Doyle C., Baxeer E.G., Hell G., Welson C.R., Miklos G.L.G.,

RA WANN K.H., Doyle C., Baxeer E.G., Hell G., Welson C.R., Miklos G.L.G.,

RA MARIA W.H., Banu A., Baxendale J. Bayraktaroglu L., Beasley R.M.,

RA HARLES W.H., Banu A., Baxendale J., Bayraktaroglu L., Beasley R.M.,

RA HARLES W.H., Banu A., Baxendale J., Bayraktaroglu L., Beasley R.M.,

RA HARLES W.H., Banu A., Baxendale J., Bayraktaroglu L., Beasley R.M.,

RA HARLES W.H., Banu B.P., Banddari D., Bolankov S.,

RA Ballev R.H., Banu A., Baxendale J., Bayraktaroglu L., Beasley R.M.,

RA HARLES W.H., Banu B.P., Banddari D., Bactler R.M., Changer R.M., Doyle R.M., Bayraktaroglu L., Beasley R.M.,

RA Ballev R.H., Banu B., Banddari B.P., Banddari J.M.,

RA Ballev R.H., Banu B., Banddari B.P., Banddari J.,

RA GLOSSE R., Gobriell J.H., Gu Z., Gaan P., Harris M.H.,

RA GLOSSE R., Harry B., Harles R.H., Rez. Rannison J.A., Ratchaman M.M.,

RA HARLES B., Bandton K.A., Hooland T.J., Hernades J.R., Bouck J.,

RA HARLES B., Bandton K.A., Hooland T.J., Well M.H., Hogsen R.M.,

RA HARLES B., Bandton K.A., Hooland T.J., Well M.H., Holland K.A.,

RA HARLES B., Bandton K.A., Hooland T.J., Well M.H., Holland K.A.,

RA HARLES B., Bandton K.A., Hooland T.J., Well M.H., Holland K.A.,

RA HARLES B., Bandton K.A., Hooland T.J., Well M.H., Holland K.A.,

RA HARLES B., Bandton K.A., Hool
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Best Local Similarity 100.0%; Pred. No. 69;
Matches 7; Conservative 0; Mismatches
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01-MAY-2000
REV PROTEIN.
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GOURFA;

GOURF
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STRAIN-SYMADESTAC;
Bear B E., Beiles E., Dapolito G., Campbell B.J., Goeken R.M.,
Bear B E., Beiles E., Dapolito G., Campbell B.J., Goeken R.M.,
Arthelm M.K., Marthum P.D., Gallo R.C., Bernard J., Zagury D.J.,
Franchini G., Sharp P.M., Hirach V.M.;
Pranchini G., Sharp P.M., Hirach V.M.;
Patterns of genopic sequence divers N.M.;
Patterns of genopic sequence divers that L. Heest monkeys (Ceropitheous liments) Are a natural leaftivitus rearroir ...

BUBLI, Arpf89115, AMPO7373, J. ...
BUBLI, Arpf89115, AMPO7373, J. ...
BUBLI, Arpf89115, AMPO7373, J. ...
BUBLI, Arpf89116, AMPO7373, J. ...

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              STRAIN-CLUMBIA.
STRAIN-CLUMBIA:
Kaneko T., Kato S., Nakamura Y., Asamisu E., Tabata S.;
Submitted (MNY-2000) to the EMBL/GenBank/DBBJ databases.
[2]
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87 IPHLAGT 93
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iruses; Retroid viruses; Retroviridae; Lentivirus
DI_TaxID=11723;
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RA WHALTE O., Easen J.A., Heidelberg J.F., Hickey E.K., Richardson D.L.,

RA Doddson R.J., Haft D.H., Govinn M.L., Nelson W.C., Richardson D.L.,

RA Manathewan J.J., Lam F., McDonald L., Utterback T., Kalewski, T.,

RA Wanathewan J.J., Lam F., McDonald L., Utterback T., Ralewski, T., Ralewski
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STRAINE-1003699; pubmed-10567266;

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A. White O., Elsen J.A., Heidelberg J.F., Hickey B.K., Peterson J.D.,

A. White O., Elsen J.A., Heidelberg J.F., Hickey B.K., Peterson J.D.,

A. Doddson R.J., Haff D.H., Govinn M.L., Nelson W.C., Richardson D.L.,

A. Doddson R.J., Hang L., Bamphile W. Crobary M., Shen M.,

A. Machtevan S. J. Lum F., McDonal M.L., Uttebber T. Falerski, B.C.,

A. Machtevan S. J. Lum F., McDonal M.L., Uttebber T. Falerski, B.C.,

A. Machtevan S. J. Lum F., McDonal M.L., Uttebber T. Falerski, B.D.,

A. Kraber C.M., Helson K.E., Saleberg S., Smith H.O., Venter J.G.,

A. Fraber C.M., Helson K.E., Saleberg S., Smith H.O., Venter J.G.,

Fradiculation R.L., Physical Research B.C.,

Fradiculation R.L., 1117(1999).

B. Science 286: 1171-1177(1999).
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098705 PRELIMINARY: PRT: 152 AA.
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090213 (TEMBUZE). 13 Created)
01-MAY-2000 (TEMBUZE). 13 Last sequence update)
01-UNY-2000 (TEMBUZE). 14, Last annotation update)
01-UNY-2000 (TEMBUZE). 14, Last annotation update)
01-UNY-2000 (TEMBUZE). 18 Last annotation update)
01-UNY-2000 (TEMBUZE). 19 Last sequence update)
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Q9YBZ3
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Best Local Similarity 100.0%; I
Matches 7; Conservative 0;
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C STRAINS-EXTERPATHOGENIC 0127:HS;

X REDLINES-95130370; PURMC4-97331;

A SCORE K.D.: Zhang H. Carlson L.K.;

D SCORE K.D.: Zhang H. Carlson L.K.;

I A. Cluster of fourteen genes from enteropathogenic Escherichia col

T aufficient for the biogenesis of a type IV pilua.;

MOI. MICROBIOL. 20:325-337(1996);

R MENTERPO. IPROULTO:

B PROSTITE: BEOGRAGO FRONCAL NTER_METHYL; UNKNOWN_1.

P PLASMICH SERVICE 183 AA; 20130 MM; IPZBFCED089720458 CRC64;

G SEQUENCE 183 AA; 20130 MM; IPZBFCED089720458 CRC64;
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Best Local Similarity 100.0%; I
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Best Local Similarity 100.0%; I
Matches 7; Conservative 0;
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098306;
01-007-2000 (TPEMBLIE). 15, Created)
01-007-2000 (TPEMBLIE). 15, Last sequence update)
01-007-2000 (TPEMBLIE). 15, Last samotation update)
LIPOPOLISACCIMATUR CORE BIOSINTHESIS.
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PROSITE: PS00454; FKBP_PPIASE_2: 1.
PROSITE: PS50059; FKBP_PPIASE_3: 1.
Isomerase.
SEQUENCE 152 AA; 16591 MM; 4COC
                                                                                                                                                                                                                                                                                                           Escherichia coli.
Plasmid EAF, and Plasmid p)
Bacteria; Proteobacteria; e
Escherichia.
NCBI_TaxID=562;
[1]
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34 ASGRARY 40
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142 VSDIVPP 148
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Bacillus/Clostridium
Bacterla; Firmicutes; Bacillus/Clostridium
Bacillus/Staphylococcus group; Bacillus.
MCBI_TaxID=86665;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7659373 PRELIMINARY; PRT; 183 AA.
059371; 96 (TERBELTEL 01, Created)
01.WOV-1996 (TERBELTEL 01, Last sequence update)
01.WOV-1996 (TERBELTEL 13) Last sequence update)
01.WOV-1996 (TERBELTEL 13) Last sequence update)
01.WOV-1996 (TERBELTEL 13) Last sequence update)
01.WOV-00ENIGO FILUS J (SPFU).
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|120 ENIKKPL 126
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STRAINHC-125 / JCH 9153;
TAkani H., Nakasone K., Takaxi Y.;
Submitted (NAR-2000) to the EMBL/GenBank/DDBJ databases.
EMBL: APO01516; BA806308.1;
SEQUENCE 165 AA; 18304 HH; 2839DF19064D2D51 CRC64;
                                                                                                                         SEQUENCE FROM N.A.
STRAIN-BAILD LASAND-EAP:
STRAIN-BAILD LASAND-EAP:
STRAIN-BAILD LASAND-EAP:
STRAIN-BAILD LASAND-EAP:
SCHEIL, Penerte JL., Ramer S.W., Bieber D., Wu C.Y., Schoolnik G.K.;
Schell, Penerte JL., Backer S.W., Bieber D., Wu C.Y., Schoolnik G.K.;
Tatteropethogenic Eachberichia coli: identification of a gene cluster
coding Cet bundle - Comming plies morphogeneais. ";
J. Bacteriol. 178:2613-2628(1996):
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SEQUENCE FROM N.A.
PLASMID=PB171;
Tobe T., Hayashi T., Han
Sasakawa C.;
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                                C.G.,
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Pred. No. 76;
O; Mismatches
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Pred. No. 82;
0; Mismatches 0; Indels
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Pred. No. 91;
0; Mismatches 0; Indels
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                                Schoolnik G.K.,
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059391
1D 059391
AC 059391
DT 01-NOV
DT 01-NOV
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DT 01-REPU
OR BEFPU
OR Escher
OC Bacter
OC Bacter
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QZZARZ

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C 059391 PRELIMINARY:
T 01-NOV-1996 (TREMBLER) 01:
T 01-NOV-1996 (TREMBLER) 11;
S BPU 1996 (TREMBLER) 11;
S BPU 1997 (TREMBLER) 11;
S BROWN 1997 (TR
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                                                                                                                                                                 Query Match 1.0%; So
Best Local Similarity 100.0%; I
Matches 7; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.0%; 1
Matches 7; Conservative 0;
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**Bacherichia coli adherence factor plasmid.**

**PRODIZIO: PROMAN_NTER_METRIC: UNENONN_1.*

**PRODIZIO: PROMAN_NTER_METRIC: UNENONN_1.*

**PRODIZIO: PROMAN_NTER_METRIC: UNENONN_1.*

**SEQUENCE 183 AA: 20148 MM; C871C0FF07F2844A CR
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01-NOV-1999 (TREMERE) 13, Last sequence update)
01-NOV-1999 (TREMERE) 13, Last annotation update)
HYPOTRETICAL 19 TAC PROTEIN APE1457.
APE1457 TAC PROTEIN APE1457.
Archers: Creatrchaeota: Desulfurococcales; Desulfurococcaceae;
Archers: Creatrchaeota: Desulfurococcales; Desulfurococcaceae;
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72 ERLQDFD 78
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NCBI_TaxID=56636;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Erwinia herbicola.
Bacteria; Proteobacteria;
Ervinia.
NCBI_TaxID=557;
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|||||||
120 ENIKKFL 126
                                                                157 VIARYGK 163
|||||||
218 VIARYGK 224
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STRAIN-EMILS.
Wright S.A.I., Beer S.V.;
Submitted (DEC-1959) to the EMBL/GenBank/DDBJ databases.
SMBHI. UB1376; AAD00513.1;
SEQUENCE 235 AA; 25733 MR; 251EC70BF1185A67 CRC64;
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9 (TrEMBLrel.
9 (TrEMBLrel.
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Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Created)
Last sequence update)
Last annotation update)
                                                                                                                                                                          Score 7; DB 2; Length 225;
Pred. No. 1.1e+02;
0; Mismatches 0; Indels
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RESULT 39
09414 PRELIMINARY: PRT; 238 AA.

AC 099414
AC 099414 PRELIMINARY: PRT; 238 AA.

AC 099416
DT 01-MAY-1997 (TERBULFAL 03, Least sequence update)
DT 01-MAY-1997 (TERBULFAL 03, Least servence update)
DT 01-MAY-1997 (TERBULFAL 06, Least annotation update)
DT 01-MAY-1998 (TERBULFAL 06, Least annotation update)
DR AAD10 PROTEIN, PARTIAL (TRANSHERY)

OC ENAMY-1998 (TERBULFAL 07, PARTIAL (TRANSHERY)

OC ENAMY-1998 (TERBULFAL 07, PARTIAL 18, PART
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S6836

AC 056836

AC 05683
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Q9KLG8
ID Q9KLG8
AC Q9KLG8
DT 01-OCT
DT 01-OCT
DT 01-OCT
DT VCAO77
OS Vibrio
OC Bacter
OX NCBLT,
RP SEQUEN
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Best Local Similarity 100.0%; Pred. No. 1.2
Matches 7; Conservative 0; Mismatches
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Best Local Similarity 100.0%; Pred. No. 1.7
Matches 7; Conservative 0; Mismatches
                                                                                              Query Match 1.0%; Score 7; DB:
Best Local Similarity 100.0%; Pred. No. 1.
Matches 7; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      056836 PRELIMINARY; PRT; 257 AA. 056836; Ort-UN-1998 (TERMILE-1.06, Created) 01-UN-1998 (TERMILE-1.06, Created) 01-UN-1998 (TERMILE-1.14, Last annotation update) 17-UN-2000 (TERMILE-1.14, Last annotation update) HYPOTHETICAL 28.5 KAN PROFIN. Wibrio cholerae 0139 fal phage.
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submitted (DRG.1998) to the EMIL/GAIDAN/DDM databases
1- HISCELLANEOUS: THERE ARE TWO GENES FOR SI IN S PONGE.
1- SIMILARITY BELONGS TO THE SAME FAMILY OF RIDGOMAL PROTEINS.
EMBL; AR1074565; CAA72256.1:
EMBL; AR107464, AM3346.1:
EMBL; AR107494, AM3346.1:
FYM: PFOOIDE: RIDGOMAL_SJAC: 1.
FYM: PFOOIDE: RIDGOMAL_SJAC: 1.
FYM: PFOOIDE: RIDGOMAL_SJAC: 1.
EMOURE: POOIDE: MIDGOMAL_SJAC: 1.
EMOURE: POOIDE: MIDGOMAL_SJAC: 1.
EMOURE: POOIDE: MIDGOMAL_SJAC: 1.
EMOURE: POOIDE: MIDGOMAL_SJAC: 1.
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OPKLAB:
OPKLAB:
OF CAME AND (TREMBLIEL: 15, Created)
OI-CCT-2000 (TREMBLIEL: 15, Last sequence update)
OI-CCT-2000 (TREMBLIEL: 15, Last annotation update)
HYPOPHERICAL PROFIEN VCA0776.
VLAUTO cholerae
Bacteria: Proteobacteria: gamma subdivision: Vibrior
KEBI_TARLD-660
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224 DAQKLLE 230
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REDLINES-943-7190; PubMed-9311128;
REDLINES-943-7190; PubMed-9311128;
REDRIA M., Shimodori S., Kojima F., Ichinose Y., Hirayama T.,
Albert H.J. Supavat K., Homma Y., Ivanaga M., Amako K.;
*Characterization of filamentous phages of Vibrio cholerae 0139 and
**Characterization of filamentous phages of Vibrio cholerae 0139 and
**Characterization of filamentous phages of Vibrio cholerae 0139 and
**Characterization of filamentous phages of Vibrio cholerae 0139 and
**Characterization of filamentous phages of Vibrio cholerae 0139 and
**Characterization of filamentous phages of Vibrio cholerae 0139 and
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**Characterization of filamentous phages of Vibrio cholerae 0139 and
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**Characterization of filamentous phages of Vibrio cholerae 0139 and
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**Characterization of filamentous phages of Vibrio cholerae 0139 and
**Characterization of filamentous phages of Vibrio cholerae 0139 and
**Characterization of filamentous phages of Vibrio cholerae 0139 and
**Characterization of filamentous phages of Vibrio cholerae 0139 and
**Characterization of filamentous phages of Vibrio cholerae 0139 and
**Characterization of filamentous phages 0139 and 0139
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24 SRLLQER 30
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MEDILINE-97051923; PubMed-8896557;
MEDILINE-97051923; PubMed-8896557;
MEDILINE-97051923; PubMed-8896557;
Medicer C., Mandel J.L., Cancel G., Abbas N., Ductr A., Didierjean O.,
Stevanin G., Agid Y., Brice A.;

"Clonain G. Kagid Y., Brice A.;

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                                                                                                                                                                                                                                                                                 Homo sapiena (Human).
Bukaryotta; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primares; Catarrhini; Hominidae; Homo.
1912_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9 11:14:16 2001
                                                                                                                                                                                                                                                                                                                                                        PERS Microbiol, Lett. 154:293-301(1997).
EMBL, D89074; BAA24163.1; -.
Hypothetical protein.
SEQUENCE 257 AA; 28548 MM; P8ZAD648273118BA CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=70203;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  238
; 26251 MW; BFBBC701DD88A2CE CRC64;
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                                                                    ; DB 12;
1.3e+02;
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AC 094438;
DT 01-MAX:
DT 01
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086430
ID 086430
AC 086430
DT 01-NOV
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AN HELDEN K. E., CLAYTON R.A., GALLS R., Gwinn M.L., Dodson R.J.,

AN HELDEN K. E., CLAYTON R.A., MILEN J.A., LINNET K.D., GATTER M.H.A.,

AN AND STEWART A.M., CONTON M.D., Pract M.S., PHILLIPS C.M., RICHARDSON D.,

AN HELDEN G. J., SHITON G.G., Fleishmann R.D., Eisen J.A., White O.,

AN HELDEN G.J., SHITON M.D., Pract M.S., PHILLIPS C.M., MILLIPS C.M., White O.,

AN HELDEN G.J., SHITON M.D., Pract M.S., PHILLIPS C.M., White O.,

AN HELDEN G.J., SHITON M.D., Pract M.S., PHILLIPS C.M., White O.,

AN HELDEN G.J., SHITON M.D., PRACT J. (2012) SHIP COLOR M. M. (2012) SHIP C.M., WHITE C.M., WHI
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Best Local Similarity 100.0%;
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Best Local Similarity 100.0%; &
Matches 7; Conservative 0;
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01-MX-1999 (TEMBLEel. 10, Last sequence update)
01-UN-2000 (TEMBLEEl. 14, Last sequence update)
40S REDSOMAL PROTEIN SN.-B (S1-B).
8831-2 OR REDS.-2 OR RED.-2 TEMBLEEL 14, Last nontition update)
8831-2 OR REDS.-2 OR SECONOMIC PROTEIN SN.-B (S1-B).
8831-2 OR REDS.-2 OR SECONOMIC PROTEIN SN.-B (S1-B).
8831-2 OR REDS.-2 OR SECONOMIC SOME STATEMENT OF STATEMENT 
086430 PRELIMINARY; PKT; 086430; 01-NOV-1998 (TrEMBLrel. 08, Created)
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203 IARYGKV 209
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099008 9 (TERMILE) 12 Created)
01-NOV-1999 (TERMILE) 13 Last sequence update)
01-UN-2000 (TERMILE) 14 Last amnotation update)
01-UN-2000 (TERMILE) 14 LAST amnotation update)
01-UN-2000 (TERMILE) 14 METHILIRANSFERASE, PUTNITE.
M1095-M1095.
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HEDILINE-2040533: PubMed1952301;
DOMBON R.J. Haft D.H. Hichey B.K., Peterson J.D., Umayam L.A.,
DOMBON R.J. Haft D.H. Hichey B.K., Peterson J.D., Umayam L.A.,
Ernolaeva H.D., Vamakhevan J., Bass S. Oin H., Drayol I. Sellers P.,
McDonald L., Ufterback T., Flainhmann R.D., Nicrman M.C., White O.,
McDonald L., Smith H.O., Colvell R. R., Mekalanos J.J., very
Fraer C.M.;
Praer C.M.;
On the Dombon Service of the cholera pathogen Vibrio
DDM, sequence of both chromosomes of the cholera pathogen Vibrio
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STRAIN=972;
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EMBL; AE004405; AAF96674.1; -.

TIGR: VCA0776; -.

SEQUENCE 246 AA; 27057 MW;
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Transferase; Ribosomal protein; Methyltransferase.
SEQUENCE 264 AA; 30280 MM; E21263892BE817AC CRC64;
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Bacteria: Thermotogales: Thermotoga.
NCBI_TaxID=2336;
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40 TVQAAAE 46
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STRAIN-972;

STRAIN-972;

Shematorova E.M., Rocol'chuk O.L., Jabedenho E.N.;

Shematorova E.M., Rocol'chuk O.L., Jabedenho E.N.;

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RA Adams M.D., Celinker S.E., Richards S., Amburner M., Henderson S.M.,

RA Gocyge R.A., Levis S.E., Richards S., Amburner M., Henderson S.M.,

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RA Golden N., Levis S.E., Richards S., Bardy M.D.,

RA Balley R.M., Basu A., Bardhale J., Bayrisktargylu L., Beasley E.M.,

RA Besson K.Y., Benso P.V., Berma B.P., Bhandari D., Baldwin D.,

RA Balley R.M., Basu A., Bardhale J., Bayrisktargylu L., Beasley E.M.,

RA Besson K.Y., Benso P.V., Berma B.P., Bhandari D., Baldwin D.,

RA Gorden K.C., Busan D.A., Buller H., Cadigu E., Center A., Chandra I.,

RA Godek N. C., Busan D.A., Buller H., Cadigu E., Center A., Chandra I.,

RA Godek N. X.J., Evangelista C.C., Perrax C., Ferrirar S., Pietschmann M.,

RA Godek N., Cong P., Gorreil J.M., Od E., Guan P., Bartis M.,

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RA Haris M.L., Harvey D., Helman T.J., Hennandes J.R., Bouck J.,
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Best Local Similarity 100.0%;
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Barbrook A.C.

Thesis (1995), University of Cambridge, C

EMEL, A0224965; CAN12258 1: -

LYTERPRO; LYRO00155; -

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Q9VKB5;
Q1-MAY-2000 (TIEMBLE-1.
Q1-MAY-2000 (TIEMBLE-1.
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082397; 98 (TREMELTAL 08, Created)
01:NOV-1998 (TREMELTAL 08), Created)
01:NOV-1998 (TREMELTAL 08), Last sequence update)
01:NOV-1998 (TREMELTAL 14, Last annotation update)
HTPOTRETICAL 3.2 6 KAN PROPELIN.
730120.2 Arabidopsis thaliana (Mouse-ear cress)
Arabidopsis thaliana (Mouse-ear cress)
Arabidopsis viridiplantae: Embryophyta: Tracheophyta: Spermatophyta:
Bukaryota; Viridiplantae: Embryophyta: Residentic Nosides: eurosids II:
MSG_TaxID=3702;
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01-COY-2000 (TREMELPA). 15 Last annotation update)
PREPROTEIN TRANSLOCASE (PRACHENT).
SEC,
Prochloror didenni.
Bacteria (Yanobacteria: Prochlorophytes: Prochloraceae:
NCD_TaxID=1216;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Drosophila melanogaster (Fruit fly),
Bukaryota; Metazos; Arthropoda; Tracheata; Hexapoda; Insecta;
Bukaryota; Meophera; Endopherysota; Diptera; Brachycera; Muscomorpha;
Ephydroides; Drosophilade; Drosophila.
MCBL_TaxID=7227;
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253 EGPECKS 259
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INTERPRO; IPRO000896; -
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SEQUENCE 288 AA; 32287 HW;
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STRAIN=CV. COLUMBIA;
Rounsley S.D., Lin X., Kaul
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01.MAY 2000 (TEDRICT: 1.3 L
01.MAY 2

Created)
Last sequence update)
Last annotation update)

PRT;

299

Sulfolobales;

Sulfolobaceae;

Sulfolobus

SEQUENCE FROM N. N.
STEALM-DOMING.

STEALM-DOMING.

MIGH spontaneous mutaion rate in the hyperthermophilic archaeon sulfolobus soliatarious is mediated by transposable elements.\*;

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RESULT 46

Q22395

PRELIMINARY: PRT: 288 AA.

D22395

D2 Q2395

D2 Q3395

D2 Q3395

D2 Q3395

D2 Q3397

D3 ANV-1996 (TEMBLICA! 01. Careted)

D7 Q1-MAY-2000 ((TEMBLICA! 01. Last sequence update)

D7 Q1-MAY-2000 (TEMBLICA! 13. Last annotation update)

D7 Q1-MAY-2000 (TEMBLICA! 13. Last annotation update)

D8 T11A5.5.

OS Cemoriabbitis elegans

OC Enkaryota, Metazoa, Mematoda; Chromadores; Rhabditida; Rhal

OC Enkaryota, Metazoa, Mematoda; Chromadores; Rhabditida; Rhal

OC Rhabditidae; Peloderinae; Caenorhabditis.

OX NCDI_TAXID=6239;

RH [1]

NCDISTA PROPER PROM N.A.

RH [1]

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Query Match 10.00; Score 7; DB 5; Length 313; Best Local Similarity 100.00; Pred. No. 1.5e+02; Matches 7; Conservative 0; Mismatches 0; Indels

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RL Submitted (JAN-2000) to the EMBL/GenBank/DDBJ databases.
DR EMBL, AZ77992; CASE5518 1. 104. Score 7: DB 1: Length 199;
Best Local Similarity 100.0N; Pred. No. 1.5e+02;
Matches 7: Conservative 0: Hismatches 0: Indels 0: Gaps 0:

OV 597 SLESAVK 593

DD 248 SLESAVK 593

DD 248 SLESAVK 593

DD 248 SLESAVK 593

DD 248 SLESAVK 593

DD 031993; PRELIMINARY; PRT; 313 AA.
AC 093893; PR
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Oy 370 RTILEAS 376

Db 91 RTILEAS 97

RESULT 50

Q04114 PRELIMINARY: PRT: 317 AA.

AC Q04114 PRELIMINARY: PRT: 317 AA.

D04114 PRELIMINARY: PRT: 317 AA.

AC Q04114 PRT: 317 AA.

AC Q04
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Search completed: April 6, 2001, 16:59:40 Job time: 234 sec

53 5 0.8 90 1 BX87_BONNO P36741 bombyx mori   54 6 0.8 90 1 BX80_BONNO P36515 bombyx mori   55 6 0.8 90 1 BX80_BONNO P36515 bombyx mori   56 0.8 90 1 BX80_BONNO P36515 bombyx mori   57 6 0.8 90 1 BX81_BONNO P3675 bombyx mori   58 0.8 90 1 BX81_BONNO P3675 bombyx mori   59 1 BXA1_BONNO P3675 bombyx mori   50 0.8 90 1 BXA1_BONNO P3675 bombyx mori   50 0.8 110 1 BXA1_BONNO P3675 morombor   50 0.8 110 1 BXA1_BON	Mon Apr 9 11:14:15 2001 us-08-403-803-2_copy_45_750.olig.rap	Copyright (c) 1993 - 2000 Compugen Ltd.  OM protein - protein search, using aw model Rum on: April 6, 2001, 16:56:42; Search time 13:55 Seponds Perfect score: 706 Registrate: 15802-403-803-2_copy_45_750 Registrate: 0.1200 Recorded: 88757 seqs, 32294092 residues Rocal state: 6 Rocal number of hits satisfying chosen parameters: 905 Hinimum DB seq length: 00 Maximum DB seq length: 2000000000 Rost-processing: Listing first 1000 summaries Database: Svissprot_39:*  Pred. No. is the number of results predicted by chance to have a soure greater than or equal to the score of the seault bullion.  Pred. No. is the number of results predicted by chance to have a soure greater than or equal to the score of the result bullion.  Result Score March Length DB ID Description  1 706 100.0 750 1 PSH_UMMAN Q04699 home septen producted by analysis of the total score distribution.  1 706 100.0 750 1 PSH_UMMAN Q04699 home septen producted by analysis of the total score distribution.  1 706 100.0 750 1 PSH_UMMAN Q04699 home septen producted prod	Non Apr 9 11:14:15 2001 us-08-403-803-2_copy_45_750.olig.rsp
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	Page 11		Page 9
\$22 6 0.8 559 1 CIRSL BRANA P27829 bradyrhizab 524 6 0.8 551 1 KITH_CAREE P1273 arian eraph 525 6 0.8 551 1 CRES_RITER P1274 P1273 arian eraph 525 6 0.8 551 1 CRES_RITER P1274 P1275 arian eraph 527 6 0.8 554 1 TRIC_RYCLE P1287 P1287 P1287 Brackbooms 6 0.8 554 1 TRIC_RYCLE P1287 P1287 P1287 Brackbooms 6 0.8 554 1 TRIC_RYCLE P1287 Brackbooms 6 0.8 554 1 TRIC_RYCLE P1287 P1287 Brackbooms 6 0.8 554 1 TRIC_RYCLE P1287 Brackbooms 7 555 1 TRIC_RYCLE P1287 P1287 Brackbooms 7 555 1 TRIC_RYCLE P1287 P1287 Brackbooms 7 555 1 TRIC_RYCLE P1287 Brackbooms 7 555 1 TRIC_RYCLE P1287 Brackbooms 7 555 1 TRIC_RYCLE P1287 P1287 Brackboom	403-803-2_copy_45_7	420 421 422 423 424 424 425 426 427 427 428 428 429 429 429 420 420 420 421 421 422 422 422 423 424 425 426 427 427 428 428 428 428 428 428 428 428 428 428	403-803-2_copy_45_7

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	Page		Page

ALIGNMENTS  RESULT 1  PSM_HIMAN STANDARD; PRT; 750 AA.  AC Q04609; D7 01-3094 (Rel. 29, Created) D7 01-3094-3994 (Rel. 29, Created) D7 01-3094-3994 (Rel. 29, Last sequence update) D8 PSW-27586 (Rel. 39, Last sequence update) D9 PSW-27586 (Rel. 39, Last sequence update) D9 PSW-27586 (Rel. 39, Last sequence update) D9 PSW-27586 (Rel. 39, Last sequence update) D8 PSW-27586 (Rel. 39, Last sequence update) D9 PSW-27586 (Rel. 39	879 6 0.8 2431 1 POLIX SFV  881 6 0.8 2447 1 LORP.MONES  882 6 0.8 2442 1 CDR_HUMAN  883 6 0.8 2442 1 CDR_HUMAN  883 6 0.8 2472 1 SPCM_HUMAN  884 6 0.8 2472 1 SPCM_HUMAN  885 6 0.8 2472 1 SPCM_HUMAN  886 6 0.8 2472 1 POLIX_ERVY  887 6 0.8 2485 1 POLIX_ERVY  888 6 0.8 2485 1 POLIX_ERVY  889 7 6 0.8 2485 1 POLIX_ERVY  889 8 6 0.8 2485 1 POLIX_ERVY  889 8 6 0.8 2481 1 POLIX_ERVY  889 8 6 0.8 2514 1 POLIX_ERVY  889 9 6 0.8 2514 1 POLIX_ERVY  890 9 6 0.8 2514 1 POLIX_ERVY  891 9 6 0.8 2514 1 POLIX_ERVY  892 1 6 0.8 2514 1 POLIX_ERVY  893 6 0.8 2514 1 POLIX_ERVY  894 6 0.8 2514 1 POLIX_ERVY  895 6 0.8 2514 1 POLIX_ERVY  896 6 0.8 2514 1 POLIX_ERVY  897 6 0.8 2514 1 POLIX_ERVY  898 6 0.8 2514 1 POLIX_ERVY  998 6 0.8 2514 1 POLIX_ERVY  998 6 0.8 3175 1 POLIX_ERVY  998 6 0.8 3175 1 POLIX_ERVY  998 6 0.8 311 1 POLIX_ERVY  998 6 0.8 311 1 POLIX_ERVY  998 6 0.8 311 1 POLIX_ERVY  999 6 0.8 311 1 POLIX_ERVY  990 6 0.8 311 1 POLIX_ERVY  900 6 0.8 311 1 PO	Mon Apr 9 11:14:15 2001 us-08-403-803-2_copy_45_750.olig.rsp	777 6 0.8 1141 MEMA_CAREL  778 6 0.8 1145 C SPAL_BACTN  779 6 0.8 1145 C SPAL_BACTN  779 6 0.8 1155 C SPAL_BACTN  779 6 0.8 1157 XYML-KNEXN  779 6 0.8 1169 C SPAL_BACTN  779 6 0.8 1170 C SPAL_BACTN  779 6 0.8 1171 C SPAL_BACTN  770 6 0.8 1172 C SPAL_BACTN  770 6 0.8 1174 C SPAL_BACTN  770 6 0.8 1175 C SPAL_BACTN  770 6 0.8 1176 C SPAL_BACTN  770 6 0.8 1177 C SPAL_BACTN  770 6 0.8 1177 C SPAL_BACTN  770 6 0.8 1178 SPAL_CARCTN  770 6 0.8 1179 C SPAL_BACTN  770 6 0.8 1170 C SPAL_BACTN  770 6 0.8 1170 C SPAL_BACTN  770 6 0.8 1171 C SPAL_BACTN  770 6 0.8 1172 C SPAL_BACTN  770 6 0.8 1174 C SPAL_BACTN  770 6 0.8 1175 C SPAL	Mon Apr 9 11:14:15 2001 us-08-403-803-2_copy_45_750.clig.rsp
		Page 19		Page 17
CARBOHYD 476 476 CARBOHYD 488 587 SINILAR 418 567 COMPLICE 750 AA: 84330 DREY MALTO 100 00 Best Local Similarity 100 00 Best Local Similarity 100 100 Matches 706: Conservative 111111111111111111111111111111111111	CC FUNCTION: MAY INTERACT WITH TRANSPERRIN OR ANOTHER LIGAND AND  CC SOSSIBLY PACILITATES METASTATIC SPERAD.  CC SUBCELLIAR LOCATION: WYPE II MEMBRANE PROTEIN.  CC TISSUE SPECIFICITY: PROSTATIC EPITHELIAL CELLS.  CC TISSUE SPECIFICITY: PROSTATIC EPITHELIAL EPITHELIAL CELLS.  CC TISSUE SPECIFICITY: PROSTATIC EPITHELIAL CELLS.  CC TISSUE	Mon Apr 9 11:14:15 2001 us-08-403-803-2_copy_45_750.olig.rsp	### 1932   1 VACL_HELPY   048258 helicobacte   0.8 1282   1 PROC_MYCGE   2012   2022	Mon Apr 9 11:14:15 2001 us-08-403-803-2_copy_45_750.cliq.rsp

QRI. VS

Page 24

Page 22

RESULT

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RESULT 8

HBA_ELEEL STANDARD: PRT; 142 AA.

LD HBA_ELEEL STANDARD: PRT; 142 AA.

AC P1450;
DT 01-JAN-1990 (Pei. 13, Created)
DT 01-JAN-1990 (Pei. 13, Last amnotation update)
DT 01-JUN-1994 (Pei. 29, Last annotation update)
DE HENGLOBIN AFRA (HEIN)

OS Electrophorus electricus (Electric eel).

OC ENHANCYDEN; MEAGAC, Chordate, Cranite; Vertebrata; Eutr
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RLIG_BACGT STANDARD; PRT; 120 AA.
110 PRIIG_BACGT STANDARD; PRT; 120 AA.
110 PRIIG_BACGT STANDARD; PRT; 120 AA.
111 PRIIG_BACGT STANDARD; PRT; 120 AA.
112 PRIIG_BACGT STANDARD; PRT; 120 AA.
113 PRIIG_BACGT STANDARD; PRT; 120 AA.
114 PRIIG_BACGT STANDARD; PRT; 120 AA.
115 PRIIG_BACGT STANDARD; PRT; 120 AA.
116 PRIIG_BACGT STANDARD; PRT; 120 AA.
117 PRIIG_BACGT STANDARD; PRT; 120 AA.
117 PRT; 120 AA.
118 PRT; 120 AA.
119 PRT; 120 AA.
110 PRT
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HSSP; P18315; 1KRC.
LWTEEDFRO, IFF002019;
FRAM, PF00659; Urease_beta: 1.
HYdrolase 124 AN; 14142 MP;
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P57047: P44559;
7 01-807 1995 (Rel. 40; Last sequence update)
7 01-607-2000 (Rel. 40; Last sequence update)
7 01-607-2000 (Rel. 40; Last sequence update)
8 SEC: INDEPENDANT PROFERIN TRANSICCASE PROTEIN TATE.
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PIR: B39102; R3858F;
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PRAM; PF00061; R1bosomal_L18p; 1.
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MEDLINE-91093287: PubMed=1985969;
RenaNrishnan V., Gerchman S.E.;
Cloning, sequencing, and overrexpression of proceins from Backlilus steatorthermophilus.*;
J. Biol. Chem. 266:880-885(1991).
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STRAIN-RB / KRO?
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HODINE-55350630; Pubmed-7542800;
Pleischmann R.D. Adams M.D. White O., Clayton R.A., Kirkness E.F.,
Fleischmann R.D. Adams M.D. White O., Clayton R.A., Kerriew J.H.
Kerlawage A.R., Bult C.J., Tomb J.-F., Desubarry B.A., Goosyne J.D.,
McReinawage A.R., Button G., Fitzhiugh W., Fields C.A., Goosyne J.D.,
McReinawage A.R., Button G., Pitzhiugh W., Fields C.A., Goosyne J.D.,
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MEDLINE=89228561; PubMed=2713104;
Huber F., Braunitzer G.;
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Inertitude produced through a collaboration the Burgean Bioinformatics and the Engli outstainon the Burgean Bioinformatics of the Commence o
                               This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swise Institute of Bioinformatics and the ZMBL contraction the European Bioinformatics Institute. There are no restrictions on its modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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STRAIN-B043/JADPI JADPI JADPI
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Bacteria; Proteobacteria; gamma
Acinetobacter.
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Transport; Protein transport; Translocation;
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Weidman J. P., Phillips C.A., Spriggs T., Hedbom E., Cotton M.D.
Utterback T.R., Hana M.C., Früger D.T., Saudek D.M., Brandon R.
Finet L.D., Prichman JL., Früger D.T., Saudek D.M., St.H.,
Goelman C.L., McDonald L.A., Small R.V., Fräser C.M., Smith H.O.,
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PRIL: 105770, AAC37155.1: -.
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INTERPRO: INTERPR
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STRAIM-S788C;

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REPUER M. Brucckner M., Schaefer M., Mueller-Auer S.;

Caser M., Brucckner M. Schaefer M., Mueller-Auer S.;

Chemosom and Ingla of 200 Kilobases from Saccharomyces

Chemosom and I. 100 (1997).
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INTERPRO: INFRO0155; .
PRAM: PRO0151 RIF-Gomain; 1.
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TIGR; MJ0461; -
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R SGD: 90044038; RR90.

R FFRO: PRO01865;

R FPAM: PF00118; RIDOGOMAL.52.1:

R FRITTS: PR00395; RIBOGOMAL.52.1: 1.

R FRITTS: PR00395; RIBOGOMAL.52.1: 1.

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STRAINS-AUCC 14827 / SROZIN
STRAINS-AUCC 14827 / SROZIN
HEDLINES-7190279; PubMed-9038161;
Norleach J. Blobberg A.;
**Merabolic and regulatory changes associated with growth of Saccharomyces cerevisiaes in 1.4 M NGC1. Evidence for compute Saccharomyces cerevisiaes in 1.4 M NGC1. Evidence for compute Saccharomyces cerevisiaes in 1.4 M NGC1 and the dihydroxyacetone pathway.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE-95203288: PubMed-7895733;
Garrela J.I., Futcher B., Kobayaani R., Latter G.I.,
VOlpe T., Marner J.R., McLaughlin C.S.;
Procein identifications for a Saccharomyces cerevis
acchase.
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RP SEQUENCE OF 52-55; 101-110 AND 134-145.

RR STRAIR-ATCC 4427; VRO2N 104

RX MEDILIES-9190279; VANDEN-9038161;

RX MEDILIES-9190279; VANDEN-9038161;

RX MEDILIES-9190279; VANDEN-9038161;

RX MEDILIES-9190279; VANDEN-9048161;

RX MEDILIES-9190279; VANDEN-9048161;

RX MEDILIES-9190279; VANDEN-9048161;

RX MEDILIES-919029; VANDEN-9048161;

RX MEDILIES-91904; VANDEN-9048161;

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CELO_CHICK STANDARD: PRT: 375 AA.
AC P19336:
DT 01-NOV-1990 (Rel. 16, Created)
DT 01-NOV-1990 (Rel. 34, Last asquence update)
DT 01-OCT-1996 (Rel. 34, Last annotation update)
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OS Gallus gallus (Chicken).
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PT DOMAIN 55 61 POLY-GLU.

PT DOMAIN 73 76 POLY-GLU.

PT DOMAIN 83 99 NOCEAR LOCALIZATION SIGNAL (POTENTIAL).

PT DOMAIN 10 10 149 HASTE LOCALIZATION SIGNAL (POTENTIAL).

PT DOMAIN 162 183 LEHOLAR LOCALIZATION SIGNAL (POTENTIAL).
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Best Local Similarity 100

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J. Bacteriol. 175:5193-5204(1993)
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SMILARITY COMPAINS I C-TERMINAL CYSTINE KNOTLIK DOMAIN CHICK.

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MEDLINE-89145205; PUNMed-2537491;

Simmons D.L.; Levy D.B.; YANDOLY, ELYMON R.L.;

Tidentification of a photol enter-repressible v-src-inducible gene.*

Proc. Natl. Acad. Sci. U.S.A. 86:1178-1182(1989).

1- FUNCTION: PROBABLE SCREETED REGULATORY PROTEIN.

1- MINICATION: BY V-SRC

1- SHILLRIFY, DELONGS TO THE INSULIN-LIKE GROWN FACTOR BINDING PROTEIN BAHLY. CEF-10/CTR61/CTF6/FISP-12/NOV PROTEIN SUBPANILY.

1- SHILLRIFY: CONTAINS 1 C-TERMINAL CYSTIRE KNOT-LIKE DOMAIN (CTCK).
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Neognathae; Galliformes; Phasianidae;
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RESULT 18
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Best Local Similarity 100.0%;
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370 IAEAVGL 376
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AR PARM, PP00425; chorismate_bind;

AR PARM, PP00425; chorismate_bind;

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30-MAY-2000 (Rel. 39, Last annotation wpdate)
MATHRANIATHE SIMPHASE COMPONENT I (EC 4-1.3.27).
TRPE OO AF1603.
Archaeglobus fulgidus.
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                                                                                   - SUBCELLIFAR LOCATION INFEGRAL MEMBRANE PROTEIN (PROBABLE).
-- SHILARTH: BELONGS TO THE NRAMP FRAILIX

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RESULT 21

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SYSTEM (18-0000005); PubMed=10844653; Maguire M.E.;

RX MEDILIRE-0000005); PubMed=10844653; B.B., Maguire M.E.;

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Best Local Similarity 100.0%;
Matches 7; Conservative (
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EMBL. Z35943; CAAS5018.1; ...
PIR: $45467; 843467.
PIR: $59224; $39224.
SCD, $5000278; VER074W.
HYPOLHELICAL PROTEIN; Hydrolase; AM
HYPOLHELICAL PROTEIN; Hydrolase; AM
HYPOLHELICAL PROTEIN; 47802 MM; 489
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STRAIN-S286;
WEDLINE-95076715; PubMed-7985423;
Van dar Aart Q.J.M., Barthe C., Doignon F., Aigle M., Crouze
Yan dar Aart Q.J.M., Barthe C., Doignon F., Aigle M., Crouze
Steensman H.V.;
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Sequence and Sequence 
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01-007-1994 (Rel. 30, Last annotation update)
01-007-1994 (Rel. 36, Last annotation update)
RYPO/HETICAL 47-8 KDA PROFIN IN HESPS-1FF32 IN
YERO/14F 00 YERO/16
Sacoharomyces cerevisiae (Baker's yeast)
Sacoharomyces cerevisiae (Baker's yeast)
Sacoharomycescae: Sacoharomycetas;
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31 SGRARYT 37
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95 LGIASGR 101
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40 71 116 144 177 216 264 305 330 337 411

1164 1164 1164 1197 1236 284 325 325 3350 3350 34 395 46652

ADJANESE.

POTENTIAL

Α,

CRC64;

0

Score 7; DB 1; ;; Pred. No. 52; 0; Mismatches

0. Length 436;

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UPTAKE

reactive

2001

us-08-403

-803-2\_copy\_45\_750.olig.rsp

Page : 469B359A66AB621C CRC64; Aminopeptidase; Zinc

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RESULT 23

NITO_METH

ID NITO_METH

ID NITO_METH

ID NITO_METH

STANDARD; PRT; 469 AA.

AC 2705.1998 (Rel. 36, Created)

DT 15-TUL-1998 (Rel. 36, Last sequence update)

DT 15-TUL-1998 (Rel. 36, Last sequence update)

DT 15-TUL-1998 (Rel. 36, Last sequence update)

DT 18-TUL-1998 (Rel. 36, Last sequence update)

DT 18
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SEAL-MODIL STANDARD; PRT; 453 AA.

AC P07039

DT 01-APR-1998 (Rel. 07, Created)

DT 01-APR-1998 (Rel. 13, Last sequence update)

DT 01-ANY-2000 (Rel. 39, Last annotation update)

DE SIGNAL RECOGNITION PARTICLE PROFIEM (FIFTY-FORM HOME)

GN FPH.

OS Bacteria; Proteobacteria; gamma subdivision; Entero Cardination and Cardin
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ce the European Bioinformatics Institute. There are no restrictions on its ce was by non-profit Institutions as long as its content is in no way ce modified and this statement is not removed. Usage by and for commorcial continues a license speciment (see http://www.isb-sib.ch/announce/ce send an email to licensesiab-sib.ch).

EMBIS 20138: 0A125871; NAT_NHT.

EMBIS 20138: 0A25871; NAT_NHT.

BR HSEP 00747; MEPSERSO, IRRO0899; .

DR HSEP 00747; PEPH DR EXCENSE EA01000; PEH STEP SO0100; REPG.

DR HSEP 500704; SREP4; 1.

DR PPAM: PPO0100; REPG.

DR PPAM: PPO0100; REPG.

DR PPAM: PPO0100; REPG.

DR PAM: PPO0100; REPG.

DR PAM: PO00100; REPG.

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Best Local Similarity 100.0%; I
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306 DIESKVD 312
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Bystrom A. S., Hyalmarsson K. J. B. Wikstroem P. M., Bjoerk G. P.;
Bystrom A. S., Hyalmarsson G. A. Besherichia coli operon containing
"The nucleotide sequence of an Besherichia coli operon containing
genes for the tRNA(mlo)methyltransferase, the ribosomal proteins $16
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86 RLGIASG 92
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; Pred. No. 53;
0; Mismatches
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RESULT 24
NIFD_METTM
ID NIFD_METTM 5
AC Q50788; P97185;
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CO - FUNCTION: NECESSARY FOR EFFICIENT EXPORT OF EXTRA-CYTOPLASHIC

CO - FUNCTION: NECESSARY FOR EFFICIENT EXPORT OF EXTRA-CYTOPLASHIC

CO - FUNCTION: NECESSARY FOR EFFICIENT EXPORT OF EXTRA-CYTOPLASHIC

CO - SUBURIT: SIGNAL RECOGNITION PARTICLE CONSISTS OF A 4.5 RNA

MOLECULE AND PROTEIN HAS A TWO DOMAIN STRUCTURE: THE G-DOMAIN

CO - ENDERGY: THE M-DOMAIN SINDS THE RNA AND ALSO BINDS THE SIGNAL

CO - SUBURIT: SIGNAL RECOGNITION FIRE OF THE RNA AND ALSO BINDS THE SIGNAL

CO - SUBURIT: SIGNAL RECOGNITION FOR THE RNA AND ALSO BINDS THE SIGNAL

CO - SINILARIT: BELONGS TO THE SEP PAMILY OF CTP-DINDING PROTEINS.

CO - SINILARIT: BELONGS TO THE SEP PAMILY OF CTP-DINDING PROTEINS.
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Best Local Similarity 100
Matches 7: Conservative
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HSSP; P00467; LMTO:
THTENPO; LP000018; ...

THTENPO; LP000018; ...

THTENPO; LP000018; ...

THTENPO; LP000018; ...

THOUSENSE_1; L1 PALSE_NEC.

PROTE; P000699; NITOOEMSE_1; L1 PALSE_NEC.

PROTE; P000699; NITOO
176 AGAKGVI 182
|||||||
51 AGAKGVI 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This SHISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMPL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to licensesiab-sib.ch).
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REDILINE-9306355; PubMade-1279430;

REDILINE-93063555; PubMade-1279430;

Signal acquence recognition by an Bacherichia coli ribonucleoprotein complex: acquence recognition by an Bacherichia coli ribonucleoprotein complex: acquence 359:741-743(1992).
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STRAINS-MIZES, STRAINS-MIZES
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MEDILINE-93063265; PubMed=1331806;
Phillips G.J., Silhavy T.J.;
The E. coli ff; gene is necessary
protein export f;
protein export f;
Rature 359:744-746(1992).
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EMBO J. 2:899-905(1983).
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genome sequence of Escherichia coli K-12.*; cience 277:1453-1474(1997).
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RA HEDITER-9616477; PubMed-9677452;

RA HEDITER-9616477; PubMed-9677452;

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RA MEDILINE-8628654; pubmed-375526;

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Best Local Similarity 100
Matches 7; Conservative
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Best Local Similarity
(4)
SEQUENCE OF 1-21 FROM N.A.
SEQUENCE OF 1-21 FROM N.A.
SEQUENCE OF 1-21 FROM N.A.
YOU J.-H., ETAULUM S.C., Hay J.G., Lemarchand P., Crystal R.G.;
YOU J.-H., EVALUEM S.C., Hay J.G., Lemarchand P., Crystal R.G.;
YULDERSHILLY Of the human airway epithelium to hyperoxia.
Constitutive expression of the oxtalase gene in human bronchial epithelial cells despite oxidant stress.
J. Clin. Invest. 93:297-302(1994).
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Hochheimer A.;
Submitted (FEB-1997) to
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MOLYBDENUM-IRON PROTEIN ALPHA CHAIN
E COMPONENT I) (DINITROGENASE).
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Pred. No.
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TRPE_HOOSH
AC P95646;
DY 15-UL-1998
DY 10-MAY-2000
DE ROCCETIA; PRE.
CO ROCCETIA
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                                                                                                           RE X BAY CRYSTALLOGRAPHY (2.75 ANGSTROMS).

RE X HEDLINE-20113173; PubMed-1066657;

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RA ADEDLINE-20113173; PubMed-10666677;

RA ADEDLINE-20113173; PubMed-10666677;

RA ADEDLINE-20113173; PubMed-10666677;

RA ADEDLINE-2011374; PubMed-10666677;

RT STUTCHUE of human erythrocyte catalase.*;

RA ACT.P. 3.65 (241.245(2000)).

CC -- FUNCTION: COCURS IN ALMOST ALL ARROBICALLY RESPIRING ORGANISMS AND SERVES TO PROTECT CELLS FROM THE TOXIC EFFECTS OF HYDROCEN

CC -- FUNCTION: COCURS IN ALMOST ALL ARROBICALLY RESPIRING ORGANISMS AND SERVES TO PROTECT ELLS FROM THE TOXIC EFFECTS OF HYDROCEN

CC -- FUNCTION: PROTECT CELLS FROM THE TOXIC EFFECTS OF HYDROCEN

CC -- SUBCLIDLAR LOCATION: PEROXISONAL.

CC -- SUBCLIDLAR LOCATION.

CC -- SUBCLIDLAR LOCATION.

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RE SEQUENCE CONTROL 1923 / 2.4.1 / NCIB 8253 / DSN 158;

RE ROADERA C. ENTO J. GIADET I.;

RE ROADERA C. ENTO J. GIADET I.;

RE SUMBLITER (FRE 159) / GIADET I.;

CC -1- CATALITIC ACTIVITY: CHORISMATE + L-GLUTAMINE = ANTHRANILATE +

CC -1- PRIVINATE + L-GLUTAMINE

CC -1- PRIVINATE = TREST STEP IN BIOSYNTHESIS OF TRYPTOPHAN.

CC -1- PRIVINATE TETRALES OF TWO COMPONENTS I AND TWO COMPONENTS II (BY

CC -1- SUBJULT, TETRALES OF TWO COMPONENTS I AND TWO COMPONENTS II PROVIDES

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15-JUL-1998 (Rel. 36, Last aequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
ANTHANNIATE SYNTHASE COMPONENT I (EC 4.1.3.27).
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BROdebacter spheeroides (Rhodopseudomonas spharroides).
Bacteria; Proteobacteria; alpha subdivision; Rhodobacter
Rhodobacter.
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RI MAI. Gen. Genet. 250:761-766(1996).

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STRAIN=168 / Sescilal:
STRAIN=95219088. Pubbe=7704263;
WEDLINE=95219088. Pubbe=7704263;
Yoshida K.-I. Sano H., Seki S. Oda M., Fuji
Yoshida K. I. Sano H., Seki S. Oda M., Fuji
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Richida K. I. Sano H. Seki S. Oda M., Fuji
Harabiology 141:337-343(1995).
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SEQUENCE FROM N.A.
STRAIN-168 / MARGURG:
MEDILINE-95189730; Pubbed-7883710;
MEDILINE-95189730; Pubbed-7883710;
MED D.G., Lindner C., Krueger S., Steinmetz M., Stuelke J.;
*New beta-9100054de [95] ogenes in Bacilius subtilis: the bglp product has both transport and regulatory functions similar to of BglF, its Sextherichia coil homolog.*;
J. Bacteriol. 177:1527-1535(1995).
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imura M., Fujita Y.; the Bacillus subtilis

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RESULT 28
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BGLP OR N17C.
Bacillus subtilis.
Bacteria; Firmicutes; Bacillus/Clostridium
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CELING D.J. Bankler A.T. Beddon A., Groenhout B.G., Nasmyth K.A.; R. STALLING D.J. Bankler A.T. Beddon A., Groenhout B.G., Nasmyth K.A.; R. Stalling D.J. 1885-434(1989).

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P08153;

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P1 01-ANG-1988 (Rel. 08, Created)

P1 01-ANG-1988 (Rel. 08, Last sequence update)

P1 01-OCT-2000 (Rel. 40, Last annotation update)

P1 01-OCT-2000 (Rel. 40, Last sequence update)

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408 GLPSIPV 414
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DY DAYS. RABIT STANDARD; PRT; 609 AA.

DY 01-AUG-1990 (Rel. 15, Created)

DY 01-AUG-1990 (Rel. 15, Last sequence update)

DY 01-AUV-1995 (Rel. 13, Last annotation update)

DZ 0XYSTROL-BINDING PROTEIN.

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OXTB_RUMAN STANDARD; PRT; 807 AA.

1D OXTB_RUMAN STANDARD; PRT; 807 AA.

1C D2559; 10 [Rel. 19; Created)

17 01-ANG-1991 (Rel. 19; Created)

17 01-ANG-1991 (Rel. 19; Last sequence update)

17 01-ANG-1991 (Rel. 13; Last sequence update)

18 OXTSTENOI-91801RG PROTEIN.

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AN COLdated, J.L.;

AN "CDRA cloning and expression of oxysterol-binding protein, an

RT oligomer with a potential leucine tipper."

BIOL Chem. 264:16796-18603(1989).

CC - FUNCTION: MAY DAY A ROLE IN THE REGULATION OF STEROL META.

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R PRESL V86895; AAC75954.1.

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G SEQUENCE: 714 AA: 77899 MF; 091289E3B7004F13 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   398 LQERGVA 404
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RC STRAIN-MAL-1 / DEM 2661 / ATCC 43067:

RX MEDILINE-96317999, PubMed-9668009?

RX MEDILINE-96317999, PubMed-9668009?

RX MEDILINE-96317999, PubMed-9668009?

RX MEDILINE-96317999, PubMed-9668009?

RX MUTCH G.J., Malte O., Olaen G.J., Zhou L., Fleischmann R.D., Goch C.J.

RA BULLT C.J., Malte O., Olaen G.J., Zhou L., Fleischmann R.D., Goch G.J.

RA BULLT C.J., Malte O., Olaen G.J., Zhou L., Fleischmann J.L., Meyven D.,

RA BULLT C.J., Malte O., Olaen G.J., Zhou R.J., Goch G.J.,

RA Gotten M.D., Roberts R.M., Malte N.A., Kadne B.P., Bordovsky M.,

RA Oction M.D., Roberts R.M., Smith H.O., Woese C.R., Venter J.C.;

RA CLEEN H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

RA R.Lank H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

RA R.Lank H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

RA R.Lank H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

RA R.Lank H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

RA R.Lank H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

RA R.Lank H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

RA R.Lank H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

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RA R.Lank H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

RA R.Lank H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

RA R.Lank H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

RA R.Lank H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

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RA R.Lank H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

RA R.Lank H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.;

RA R.Lank H.-P., Fra
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Best Local Similarity 100.0%; 1
Matches 7; Conservative 0;
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30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last sametation update)
16-MAY-2000 (Rel. 39, Last annotation update)
16-MAY-2000 (Rel. 39, Last annotation update)
17-MANSPERASE; (Rel. 39, Last annotation update)
18-MAY-2000 (Rel. 39, Last annotation update)
18-MAY-2000 (Rel. 39, Last sametation update)
18-MAY-2000 (Rel. 39, Last sametation update)
18-MAY-2000 (Rel. 39, Last sametation update)
18-MAY-2000 (Rel. 39, Last sequence u
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BEBL. ALO3397: CAM19377.1; -
INTERPRO: IPRO02819:
PRAM: PP01966 UB-chickly transferase:
PRAM: PP01966 UB-chickly transferase:
SEQUENCE 635 AA: 85046 MH; 551296D8BE77DE5D
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This SHIGAPPOT entry is copyright. It is produced through a collaboration between the System Numerical of Bioinformatics and the BMDL outstation between the Direction and the SMDL outstation of the BMD proposed Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no value by non-profit institutions as long as its content is in the second of this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.lab-sib.ch/announce/or send on cashi to licenseiab-sib.ch).
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5 SSWRGSL 561
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0; Pred. No. 92;
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INTERPRO; IPRO02203; -..

DR PPAM: PP00009; CTP_ESTU; 1.

DR PROSITE: PS000801; PROFIES SPLICING; 1.

DR PROSITE: PS000801; PROFIES SPLICING; 1.

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RN HEDIANDES FROM N.A.

RC STRAIN-HEJ37;

RA MEDIANDES 54,7138;

FOUNDATION FOR A R. R. Holden D. N.;

RI chilin synthases with different hunctions;

RT chilin synthases with different hunctions in cookers is

RT chilin synthases with different hunctions with the centre of the 
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REPALMETJA STANDARD; PRT; 1341 AA.

10 REPALMETJA STANDARD; PRT; 1341 AA.

11 SHOCKLESS (REL. 37, Last anguence update)

12 JOHN JOHN (REL. 37, Last anguence update)

13 OHAY 2000 (REL. 39, Last annotation update)

14 DE DNA-DIRECTED RRA POLYMERASE SUBUNIT A' (RE 2.7.7.6).

15 DE DNA-DIRECTED RRA POLYMERASE SUBUNIT A' (RE 2.7.7.6).

16 SHETHARDOOCCUS BLYSACHAROCTA; Methanococcases:

17 Archaes Euryachaeota; Methanococcases:

18 Methanococcus.
in SigUpmers From M. A.

STEPRINH-JALT. / DBW J661 / ATCC 43067;

RETRIANS-461 / DBW J661 / ATCC 43067;

RETRIANS-461 / DBW J681 R. J. Zhou L., Fleischmann R.D.,

RA Bult C.J., Mhite O., Olsen G.J., Zhou L., Fleischmann R.D.,

RA Sutton G.G., Blake J.A., Fliegerald L.H., Clayfon R.A., Goeayne J.D.,

RA Stribvage R., Doughter B., Weinstock K.G., Merrick J.H., Clodek A.,

RA Coverbeck R., Kirkness E.F., Weinstock K.G., Merrick J.H., Clodek A.,

RA Coverbeck R., Kirkness E.F., Weinstock K.G., Merrick J.H., Nguyen D.,

RA Cotton H.D., Roberts R.H., Hurst N.A., Maine B.P., Berodowsky H.,

RA Clotton M.D., Roberts R.H., Hurst N.A., Maine B.P., Berodowsky H.,

RA Clotton M.D., Roberts R.H., Hurst N.A., Walne B.P., Berodowsky H.,

RA Clotton M.D., Roberts R.H., Hurst N.A., Walne B.P., Berodowsky H.,

RA Clomplete Sponse sequence of the methanogenia archaeon, Methanococcus

RET Complete Sponse sequence of the methanogenia archaeon, Methanococcus

RET Science 2731058-1073(1996).
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Matches 7; Conservative
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Best Local Similarity 100.0%; I
Matches 7; Conservative 0;
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|||||||
126 KAENIKK 132
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18-2, 1ST BART (POTENTIAL),

HOAD 18-2 INTEIN (POTENTIAL),

PROBABLE TRANSLATION INITIATION PA

18-2, 2ND PART (POTENTIAL),

GTP (BY SIMILARIY),

GTP (BY SIMILARIY),

59 MM; 663PR 3587EEDDJ GRC64;
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O; Mismatches
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Pred. No. 1.2
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RX MEDLINE-91123227, PubMed=1704006;

RA ADD H. Matsumara T., Kodama T., Ohta H., Fukui R., Kato R.,

Rapawai, T., Radamara T., Kodama T., Ohta H., Fukui R., Kato R.,

RT **Agawai, T. **A
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DR INTERPRO; IPROCAT9; C. BINTAIG.1 19.
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Knochs T., Hillief L., Jier M., Johnston L., Langston Y.,
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Vignati D., Wilcox L., Wohldman P., Materaton R., Wilson R.,
Vaudin M., Rifkin L., Riibet L., SE Peter H., Trevaskis E., Vanghan K.,
Vignati D., Wilcox L., Wohldman P., Materaton R., Wilson R.,
Vaudin M.;

ence 265:2077-2082(1994).

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coding sequences of 40 new genes (KIAA0041-KIAA0080) deduced years of cDNs clones from human cell line KG-1.";
Res. 1:233-226(1994)
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EMBL; D38548; BAA07551.1; -. Hypothetical protein. SEQUENCE 1698 AA; 191188 MW;

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CHARACTERIZATION

HEDLINE-8813/S50: Dubbed-2815/S8;
HEDLINE-8813/S50: Dubbed-2815/S8;
Jackson S.P., Loasty, M., Beggs J.D.;
Jackson S.P., Loasty, M., Beggs J.D.;
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RESULT 42
PROB\_YEAST
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01-PB3-1994 (Rel. 28, Last asquence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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EMBL, 22472; CAA60854.1: .

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Shea J.E., Toya J.H., Johnston L.H.;

Shea J.E., Toya J.H., Johnston E.H.;

The budding yeast U5 snRMP Prp8 is a highly conserved

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Nucleic Acids Res. 22:5555-5564(1994).

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OS PLARMOCIUM ELICIPATUM (1801SEE CDC / HONGULES).

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AD HID\_PRESS P. STANDARD: PRT: 37 AA.
DR 13-AUG\_1897 (Rel. 05, Created)
DR 13-AUG\_1897 (Rel. 05, Last sequence update)
DR 13-AUG\_1896 (Rel. 34, Last snotation update)
DR HFO-INOSITOL-BINDIM PROTEIN (FRACHENT).
OS Pseudomonas sp.

Page 74

Bacteria;

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INTERPRO: IPRO01938; ...
PRAM: PF00314; thaumatin: 1.
PROSITE: PS00316; THAUMATIN; P
PUNGICIAE.
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44
SEQUENCE 44 AA; 4453 MW; 9

920713BCE551C1E4 PARTIAL.

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RESURD 47

HT SYMPJ

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AC P3031;
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THER MORNY
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TO STANDARD: PRT: 44 AA.

AC P3304;
PT 01-0CT-1993 (Rel. 27, Created)
DT 01-0CT-1993 (Rel. 27, Created)
DT 01-0CT-1993 (Rel. 27, Last sequence update)
DT 01-0CT-1993 (Rel. 28, Last annotation update)
DT 01-0CT-1993 (Rel. 28, Last annotation update)
DT 01-0CT-1994 (Rel. 29, Last annotation update)
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                                                    RE SEQUENCE FROM N.A.

RE MUCKLE J. W. Norty A.P., Turner J.S., Robinson N.J.;

RE MUCKLE J. W. Norty A.P., Turner J.S., Robinson N.J.;

RE MUCKLE J. W. Norty A.P., Turner J.S., Robinson N.J.;

RE MUCKLE J. W. Norty A.P., Turner J.S., Robinson N.J.;

RE MUCKLE J. W. Norty A.P., Turner J.S., Robinson N.J.;

RE MARSCHIPTIONAL CONTROL BY Green metal locus and analysis of RI Laboration of a procharyotic metaloriton in the March 1607014;

RE MARSCHIPTIONAL CONTROL BY THE ROBINSON N.J.;

RE CUBRICITESTION, SEQUENCE FROM N.A. AND PARTIAL SEQUENCE.

RE MEDIUME-92299082; PubMed-1607014;

RE CUBRICITESTION, PUBMED-1607014;

RE CUBRICITESTION N. P. HUCKLE J. W. Norty A.P., Robinson N.J.;

RE "Cyanobacterial metalothionesin gene expressed in Emberichia coli.

RT "Ketal-binding properties of the expressed protein."

RE "Cyanobacterial metalothionesin gene expressed in Emberichia coli.

RT HETAL HIGH STATE OF HUCKLE J. W. NORTH METAL ION HOMEOSTASIS

RE "Cyanobacterial metalothionesin gene expressed in Emberichia Robinson general genera
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PubMed-6735978;

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17 VDPSKA 22
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STEALINGY. MONT RISO 1508.
STEALINGY. MONT RISO 1508.
MEDLINE-90017994, PubMed-195240;
Hojsard J. Jacobson S., Svendsen I.;
"Two antifungal chanactistike protectins from barley grain.";
PERS Lact. 20112111(1991).
TRENDERS VIRTUMANAL ACTIVITY. INHIBITS THE GROWTH OF
TRENDERS VIRTUMANAL COLVETTY. INHIBITS THE GROWTH OF
THE SHICHART: BELONGS TO THE THANNATIN FAMILY.
PIR. SITS 3: 
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|||||||
28 KSYPDG 33
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oteria; Cyanobacteria; Chrococccales; Synechococcus.
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RESULT 48

OPPALABATE

ID GYPALABATE

OFFICIAL 16, Created)

DT GYPALABATE

DT GYPALABATE

DT 01-MAR-1990 [Rel. 10, Created)

DT 01-MAR-1990 [Rel. 10, Last sequence update)

DT 01-MAR-1990 [Rel. 36, Last annotation update)

DT 01-MAR-1990 [Rel. 36, Last annotation update)

DE GAS VESICLE PROFIEN (GVP).

GN GYPA.

GO Anabena flos-aquae

BACTETIA: Cyanobacteria; Nostocales; Nostocaceae:

BACTETIA: Cyanobacteria; Nostocales; Nostocaceae:

BACTETIA: Cyanobacteria; Nostocales; Description of the STRAINGCAR PROFIES (AVP).

RESTULNES FROM N. A. AND SEQUENCE OF 1-12.

RESTULNES-230109; PubMed-1527466;

RA GRIFFITHA A. F. Halby A. E. Hayes P. K.

RESTULNES-2310769; PubMed-152746;

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ITD_SYMES PROPER STANDARD; PRT; 53 AA.

AC P00003:8 (Rel. 08, Created)

DT 01:AUG-1988 (Rel. 08, Last acquence update)

DT 01:AUG-1988 (Rel. 15; Last annotation update)

DE METRICOPTION (REL. 15; Last annotation update)

DE METRICOPTION (REL. 15; Last annotation update)

OC Bacteria: Cyanobacteria: Chroococcales; Synechococcus.

RE 101

RE 2012BCC CYANOBACTERIA: Chroococcales; Synechococcus.

RE 2012BCC P1.24.

RE 2012BCC P1.25.

RE 2012BCC P1.
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Best Local Similarity 100.0%; I
Matches 6; Conservative 0;
                                                                                                                  SENGURNCE FROM N. A.
STRAIN-CCLP 1003/11F;
MEDLINE-9717655; PubMed-9063646;
Kloman R., Rayse P.K.;
"Genes encoding proteins homologous to halobacterial & L are located downstream of gypC in the cyanobacter flos aquee."
[10s aquee."]
[10s aquee."]
[11]
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17 VDPSKA 22
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STRAIN-TM. 20;
STRAIN-TM. 
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Pacteria; Cyanobacteria; Nostocales; Nostocaceae;
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Pred. No. 88;
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DR ENGL. X06005. CAA23468 1:

DR PIR. D21951. WAX2488 1:

DR PIR. D21951. WAX2488 1:

DR PIR. D21951. WAX2488 1:

DR PIR. D21951. AX7475.

NR PIR. D21951. AX7475.

NR PPAM: PPO000630;

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REDITER-0707543; PubMed-0908234;

A Mayes P.K. Walsby, A.E., Walker J.E.,

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Best Local Similarity 100.0%;
Matches 6; Conservative
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RESULT 3 TREAL Species Melanoplus anguinipes entomopoxvirus C.Species Melanoplus anguinipes entomopoxvirus C.B. Kundo, C.L.; Talian, E.R.; Lu, Z.; Oma, E.; Kutish, G.F.; Rock, D.L. S. Kundo, C.L.; Talian, E.R.; Lu, Z.; Oma, E.; Kutish, G.F.; Rock, D.L. A. Melanoplus anguinipes entomopoxvirus A. Melanoplus anguinipes entomopoxvirus A. Melanoplus anguinipes entomopoxvirus A. Melanoplus Entomopolus anguinipes entomopoxvirus A. Melanoplus (20484, MUID:980861) A. Melanoplus (20484, MUID:980861) A. Melanoplus (20484, MUID:980862) A. Melanoplus (20484, MUID:980862) A. Melanoplus (20484) A. Melanoplus (20	RESULT 2  Peptidase-like protein - Arabidopsis thaliana N.Alternate names: protein T5923.80  C:Species: Arabidopsis thaliana (mouse-car cress) C:Species: Arabidopsis thaliana (mouse-car cress) C:Compasion: 74831 R:Obermaler: B: Ottenwaelder: B: Duchemin, D: Zeitler, K.: Mewes, H.W.: Lemcke, K.: Mayer, K.F.X.: Quetier, F.: Salanou R:Obermaler: B: Ottenwaelder: B: Duchemin, D.: Zeitler, K.: Mewes, H.W.: Lemcke, K.: Mayer, K.F.X.: Quetier, F.: Salanou Arabidopsis (Compasion of the Protein Sequence Database, March 2000 A. Marches (Compasion of the Protein Sequence Database, March 2000 A. Marches (Compasion of the Protein Sequence Database, March 2000 A. Marches (Compasion of the Protein Sequence Database, March 2000 A. Marches (Compasion of the Protein Sequence Database, March 2000 A. Marches (Compasion of the Protein Sequence Database, March 2000 A. Marches (Compasion of the Protein Sequence Database, March 2000 A. Marches (Compasion of the Protein Sequence Database, March 2000 A. Marches (Compasion of the Protein Sequence Database, March 2000 A. Marches (Compasion of the Protein Sequence Database, March 2000 A. Marches (Compasion of the Protein Sequence Database, March 2000 A. Marches (Compasion of the Protein Sequence Database, March 2000 A. Marches (Compasion of the Protein Sequence Database, March 2000 A. Marches (Compasion of the Protein Sequence Database, March 2000 A. Marches (Compasion of the Protein Sequence Compasion of the Protein Seque	Mon Agr 9 11:14:14 2001 us-08-403-803-2_copy_45_750.olig.rpr Page 23	977 6 0.8 464 2 H82011  978 6 0.8 464 2 F755121  979 6 0.8 464 2 F755121  979 6 0.8 464 2 F755121  970 6 0.8 464 2 F755121  971 6 0.8 464 2 F755121  972 6 0.8 464 2 F755121  973 6 0.8 464 2 F755121  974 6 0.8 464 2 F755121  975 6 0.8 465 2 F75512  977 6 0.8 465 2 F75512  978 6 0.8 465 2 F75512  979 6 0.8 465 2 F75512  970 6 0.8 472 2 F75752  970 6 0.8 472 2 F75752	Mon Apr 9 11:14:14 2001 us-08-403-803-2_copy_45_750.olig.rpr Page 21
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A:Molecule type: m8NA
A:Residues: 1-165 COPTD-
A:Cross-reforences: pBN:AL110260
A:Experimental source: adult uterus; clone DRF2p586B1621
A:Note: DRF2p596B1621.1
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Status: preliminary
Cresides: 1-26 (ARN)
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Map Position: 3
Introne: 31/3; 74/2; 124/3; 161/2; 189/2; 205/3; 236/2; 278/3; Note: PRIZ1_210
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A.Gene: modA
A.Genome: plasmid pAO1
C.Guperfamily: molybdate-binding periplasmic protein
C.Keywords: molybdenum transport
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A.Mon Position: X
A.Mon Position: 40/1: 145/1: 230/1: 275/3: 313/1: 344/2: 405/1: 453/2: 576/2: 671/2: 713/3
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LiReference number: 222860
LiReference number: 122860
LiReference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ypothetical protein R57.1 - Caenorhabditis elegans
species: Caenorhabditis elegans
species: 13-oct-1999 sequence_revision 15-oct-1999 stext_change 18-Peb-2000
Accession: T30154
Accession: T30154
Big Bat Albrary, February 1997
Description: The sequence of C. elegans cosmid R57.
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Best Local Similarity 100.0%;
Matches 8; Conservative (
                                                                                                                                                                                                                                                 Status, preliminary, translated from GB/EMBL/DDBJ
Molecule type, DBA
Residues, 1-75, CFN>
Cross, reference, BMSL,U88179; FIDN:AAB52660.1; GBPDB:GN00028; CESP:R57.1
Experimental Source: strain Bristol N2: Clone R57
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53 FTEIASKF 60
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Pred. No. 5.2;
0; Mismatches 0; Indels
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A:Accession: A48773
A:Rolecule type: DNA
A:Residues: 1-1202 CND2>
A:Cross-references: EMBL:L06209; NID:g172441; PIDN:AAA0316B.1; PID:g172442
A:Cross-references: EMBL:L06209; NID:g172441; PIDN:AAA0316B.1; PID:g172442
A:Rote: parts of this sequence, including the amino end of the mature protein, were
R:Gentles, S.; Bowman, S.
submitted to the EMBL: Data Library. November 1994
A:Reference number: \$49827
A:Accession: \$49836
A:Rotecule type: DNA
A:Residues: 1-1202 CGNN
A:Reference number: \$61393; NUID:93028566
A:Rotecssion: \$61340 Mon A.Map position: 13L A.Genome: nuclear A.Gerome: nuclear C.Reywords: hydrolase; mitochondrion; RNA binding; tRNA processing F:123-1202/Product: ribonuclease P, mitochondrial fatatus experimental Αpr Genetids: SGD:RPM2
SGCos:references: SGD:S0004556; MIPS:YML091c :Reference number: S1339; MIID:33028566
:Accession: S6134
:Kolecule type: DNA
:Kolecule type: DNA
:Kolecule type: DNA
:Kolecule type: S1339
:Kolecule type Query Match 1.1%; Best Local Similarity 100.0%; Matches 8; Conservative 9 11:14:14 2001 0 Score 8; DB 2; Pred. No. 21; 0; Mismatches us-08-403-803-2\_copy\_45\_750.olig 0, Length 1202; Indels PID:g172442 mature protein, were 0 Gaps 0 ρĄ Page 28

RESULT 11

Places/Itansferase - Streptococcus salivatius

Cispecies Streptococcus salivatius

Cispecies Streptococcus salivatius

Cispecies 27-Oct-1999 sequence\_revision 22-Oct-1999 stext\_change 22-Oct-1999

CiAccession 19088

Risimpon, C.L.; Giffard, P.M.; Jacques, N.A.

Infect Imman, 63, 609-611, 1998

A:Title: Streptococcus salivatius ATCC 25975 possesses at least two genes oc 453 SKYDPSKA 460 673 SKVDPSKA 680

least two genes coding for primer-independent glucosyltransferase

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Ribatomate names: eukaryotic ribasomal protein protein L29 homolog

Ribatomate names: eukaryotic ribasomal protein protein L29 homolog

Ribatomate names: eukaryotic ribasomal protein protein L29 homolog

Cibatomate names: eukaryotic ribasomateriophicum

Cibatomaterio protein C5920

Ribatomaterio C5920

Riba
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T39580
UP1319 - fission yeast (Schizosaccharomyceé pombe)
C:Species Schizosaccharomyces pombe
C:Species On-Dec:1999 #sequence_revision 03-Dec-1999 #text_change 21-Jan-2000
C:Accession: 193880
RVOLKMert, G: Mood, V: Rejandream, M.A.: Barrell, B.G.
RVOLKMert, G: Mood, V: Rejandream, M.A.: Barrell, B.G.
RVOLKMert G: the EMBL Data Library, August 1997
A: Reference number: 821865
A: Recent consumer: 821865
A: Accession: 193580
A: Residues: 1-102 - VOLX
A: Cross-references: EMBL:899759, PIDN:CAB16898.1; GSPDB:GN00067; SPDB:SPBC1689.06c
A: Experimental source: strain 972h-; cosmid c1689
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A:Gene: uvill; SPDB:SPBC16E9.06c
A:Map position: 2
C:Superfamily: bolA protein
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                                                                                                                                                                           ribosomal protein L18 - Bacillus stearothermophilus
C.Species: Bacillus stearothermophilus
C.Species: Bacillus stearothermophilus
C.Species: Bacillus stearothermophilus
T.Remealon_Bapato1_19885
R.Khura, J: Manray, M.
RES.Latt. 210, 85-96, 1987
RES.Latt. 210, 85-96, 1987
N.TILE: The complete anniopti. A23102_K0TI: 97105336
A.McCession: B23102
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A:Accession: T30838
A:Status: prelianiaxy; translated from GB/EMRI/DDBJ
A:Status: prelianiaxy; translated from GB/EMRI/DDBJ
A:Kolecule type: DNA
A:Rosidues: 1-1577 GID
A:Cross-references: EMBL:L35928; NID:g662380; PID:g662381; PIDN:AAC41413.1
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Best Local Similarity 100.0%; Pred. No. 28;
Matches 8; Conservative 0; Mismatches
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1091 AFLDELKA 1098
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60 SPEFSGM 66
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21 LDELKAE 27
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A:Molecule type: DNA
A:Residues: 1.13 (MHZ)
A:Cross-references: GB:AR001836; N.
A:Experimental source: strain R1
C/Genetics:
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conserved hypothetical protein - Deinococcus radiodurans (strain R1)
cospecias: Deinococcus radiodurans
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A;Reference number: A39085; MOID:91093287
A;Recession: D93085
A;Roiceule type: DNM.
A;Residues: 9:111 < RNM.
A;Residues: 9:111 < RNM.
A;Cross:references;GB:M7624
C;Reywords: protein blosynthesis; ribosome
                                                                                                                                                                                                                                                                                                                                                            A;Gene: DRB0052
A;Map position: megaplasmid
A;Genome: plasmid
A;Note: plasmid MP1
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Best Local Similarity 100.0%; I
Matches 7; Conservative 0;
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694 TVQAAAE 700
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118 TVQAAAE 124
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11 APGVKSY 17
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61 KEFGLDS 67
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pred. No. 32;
0; Mismatches
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hypothetical protein - Delnococous radiodurans
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:Status: Pre: DIA.;
|Residus: 196 (MH1).
|Cross:references: GB AE001910; GB:AE000513; NID:96458198; PIDN:AAF10058.1; PID:96458209; TIGR:DR0518; GSPDB:GN00077
|Experimental source: strain Ri
Query Match 1.0%; Score 7; DB 2;
Best Local Similarity 100.0%; Pred. No. 23;
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Species: Deinococcus radiodurans
'Species: Deinococcus radiodurans
'Species: Deinococcus radiodurans
'Charle: O.: Dec. 1999 sequence_revision O3-Dec.1999 $text_change O9-Jun-2000
'Rocession: B75347
'Robert C. R. Stean, J. A.; Heidelberg, J.F.; Hickey, E.K.; Peterson, J.D.; Dodson, R.J.; Haft, D.H.; Gwinn, M.L.; Nelson, J. R.; Shen, M.; Vanathevan, J.J.; Lam, P.; McDonald, L.; Dtterback, T.; Zalewski, C.; Makarova, K.S.; Aravind, L.; Daly, Inclaim C. Stean, C. R. Stean, C. R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Typothetical protein PAOB59 [imported] - Pseudomonas aeruginosa (strain PAO1)
;Species: Pseudomonas aeruginosa
;Species: 15-5ep-2000 saequence_revision 15-5ep-2000 stext_change 15-5ep-2000
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Best Local Similarity 100.0%; I
Matches 7; Conservative 0;
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|120 ENIKKFL 126
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Lecule (vps: DNA)
Siduem: 1-200 (STS)
Descripterences: G:SE004520; GB:AE004091; NID:g9946745; PIDN:AAG04248.1; GSPDB:GN00131; PASP:PA0859
Peripental Source: Gttain PAO1
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C;Superfamily: BKBP-type peptidylprolyl isomerase: BKBP-type peptidylprolyl isomerase F;64-111/Domain: BKBP-type peptidylprolyl isomerase homology <PPI>
       bigo protein - Becherichia coli
C:Specie: Escherichia coli
C:pate: 11-Feb-1998 feequence_revision 20-Feb-1998 #text_change
C:Accession: 57997
R:Stone, K.D.: Ekney 5.: Carlson, L.K.: Donnenberg, M.S.
M:Stone, K.D.: Ekney 325-337, 3958
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A.Molecule type DNA
A.Reakdus; 11-73 <KAN-
A.Gross-references: DBBJAP000061; NID:95104821; PIDN:BAA80455.1; PID:d1044241; PID:95105141
A.Experimental source: strain K1
C.Genetics:
C.Genetics:
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Popecies: Aeropyrum pernix
Pichte: 20-Aug-1999 sequence_revision 20-Aug-1999 stoxt_change 20-Aug-1999
Pichte: 20-Aug-1999 sequence_revision 20-Aug-1999 stoxt_change 20-Aug-1999
Pichte: 20-Aug-1999 sequence_revision 20-Aug-1999 stoxt_change 20-Aug-1999
Pichte: 20-Aug-1999 sequence, revision 20-Aug-1999
Pichte: 20-Aug-1999 sequence of an aerobic hyper-thermophilic Crenarchaeon, Aeropyrum pernix Kl.
Pichternoenumber ynows equence of an aerobic hyper-thermophilic Crenarchaeon, Aeropyrum pernix Kl.
Pichternoenumber ynows equence of an aerobic hyper-thermophilic Crenarchaeon, Aeropyrum pernix Kl.
Pichternoenumber ynows equence of an aerobic hyper-thermophilic Crenarchaeon, Aeropyrum pernix Kl.
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Pichternoenumber (20-Aug-1999)
Pichternoenumb
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1.0%; Score 7; DB 2;
Best Local Similarity 100.0%; Pred. No. 36;
Matches 7; Conservative 0; Mismatches
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Best Local Similarity 100.0%;
Matches 7; Conservative
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34 ASGRARY 40
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|||||||
72 ERLQDFD 78
Zhang, H.Z.; Carlson, L.K.; Donnenberg, M.S.
.. 20, 325-337, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 152;
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36

A: Map position: FOR411301-411927 C: Superfamily: rat ribosomal protein S3

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Query Match 1.0%; S
Best Local Similarity 100.0%;
Matches 7; Conservative 0;
176 AGAKGVI 182
|||||||
122 AGAKGVI 128
                                          score 7; DB 2;
b; Pred. No. 48;
0; Mismatches
                                             0
                                                              Length 208
                                             Indels
                                             0
                                             Gaps
                                             0
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nitrogenase (EC 1.18.6.1) molybdenum-iron protein alpha chain - Methanobacterium therm
N.Alternate names: nitrogenase chain D [misidentification]
C:Species. Methanobacterium thermoaucorrophicum
C:Date: 28-0cf-1996 sequence\_revision 13-Mar-1997 \*\*text\_change 04-Peb-2000
C:Accession: 563551; \$57.44
R:Hochielmer, A.; Schmitz, R.A.; Thaner, R.K.; Hedderich, R.
Burr, J. Biochem 214, 910-90, 1995
A.Title: The tungsten formylmethanofuran dehydrogenase from Methanobacterium thermoauth
A.Mocession: 563515; MVID:96183477
A.Mocession: 563515; MVID:96183477
A.Mocession: 563553
A.Title: Preliminary: nucleic acid sequence not shown: translation not shown
A.Molecule type: DNA
A.Mocession: 563504
A.Gross-references: BMDL:M97971
A.Mocession: EMBL:M97971
A.Mocession: EMBL:M97971 thermosutotrophicum thermoautotrophicum CONTAINS (strain Marburg)

motifa

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RESULT 28

604 11 Docomal Protein Slac (S1). fission yeast (Schizosaccharomyces pombe)

604 12 Docomal Protein She (S1). fission yeast (Schizosaccharomyces pombe)

8. Alternate and septembers (Species Schiicosaccharomyces pombe)

9. Species Schiicosaccharomyces pombe

9. C. Accession: 56441; 73639

8. Rodell. (S. Bowan, S. S644)

8. Rodell. (S. Bowan, S. S644)

8. Rodession: 56441

8.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Procession: C. Stall Sections of Genes encoding Acinetobacter calcoaceticus procession: C. Stall Section 10-Sep-1999 stext_change 03-Mar-2000
Ribatnett, C.; Medde, E.L.: Medd
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C.Species: V.Huro cholerae
C.Species: V.Huro cholerae
C.Species: V.Huro cholerae
C.Species: V.Huro cholerae
C.Species: 18 - Aug-2000 #sequence_revision 20 - Aug-2000 #text_change 15 - Sep-2000
C.Nocession: 1892.19
R.Heidelberg, J.F.; Eisen, J.A.; Nelson, W.C.; Clayton, R.A.; Gwinn, M.L.; Dodson, R.J.; Haft, D.H.; Hickey, E.K.; Peterson
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C;Superfamily: dinitrogenase alpha chain; nitrogenase
C;Keywords: oxidoreductase
       Query Match 10%; Score 7; DB 2; Length 252; Best Local Similarity 100.0%; Pred. No. 58; Matches 7; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                            Gene: rpsl·l
;Map position: IL
;Superfamily: rat ribosomal protein S3a
;Keywords: protein biosynthesis; ribosome
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170%; Score 7; DB 1;
Best Local Similarity 100.0%; Pred. No. 55;
Matches 7; Conservative 0; Mismatches
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Best Local Similarity 100.0%; P
Matches 7; Conservative 0;
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| 240 TEWAEEN 246
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41 AETLSEV 47
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51 AGAKGVI 57
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; Pred. No. 51;
0; Mismatches
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La R. Mattire 406, 477-883, 2000

Artitle DMS, Sequence of both chromasmas of the cholera pathogen Vibrio cholerae.

A. Reference maker: 83035; MUD-20406833

A. Reference maker: 1246 cHED:
A. Residues 1-246 cHED:
A. Residues 1-256 cHED:
A. Res
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QY 257 DAGKLE 263

Db 224 DAGKLE 230

RESULT 29

T38213

4 Bribosomal protein SiB - fission yeast (Schizosaccharomyces pombe)
C:Species: Schizosaccharomyces pombe
C:Species: Schizosaccharomyces pombe
C:Astes O:Dec:1399 sequence\_revision O:Dec-1999 stext\_change O:Sep-2000
C:Astes O:Dec:1399 sequence\_revision O:Dec-1999 stext\_change O:Sep-2000
C:Astes O:Dec:1391 sequence\_revision O:Dec-1999 stext\_change O:Sep-2000
C:Astes O:Dec:1392 sequence\_revision O:Dec-1999 stext\_change O:Sep-2000
C:Astes O:Dec:1392 sequence\_revision O:Dec-1999 stext\_change O:Sep-2000
C:Astes O:Dec:1392 sequence\_revision O:Dec-1999 stext\_change O:Sep-2000
R:BaGcock, K: Oburche: 22179
A.Rocession: T3213
A.Rocession: T3213
A.Rocession: T3213
A.Rocession: T5129
A.Rocession: T512

## Overy Match Bast Local Similarity 100.0%; Pred. No. 50; Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0; 9 257 DANGLE 263 Oy 257 DANGLE 263 Lillill Db 224 DANGLE 230 RESULT 30 Species: 30-Aver-195 sequence\_revision 23-Peb-1996 stext\_change 18-Aug-2000 C.Daccession: 859364, 561622; 864876

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R.Deminnova, M.A., Permone, T.G.; Ellis, S.R.
subsitted to the 2MB Data Library, August 1995
A.Reference number: 659364
A.Moceasion: 559364
A.Moceasion: 559364
A.Molecular type: DNA
A.Moceasion: 4525 - CDBL
A.Moceasion: 651622
A.Molecular type: DNA
A.Molecular type: DNA obiquie type: DNA esidues: 1-252 (AND) ross-references: EMBL:273220; MIPS:KLRO48w; NID:91360382; PIDM:CAA97578.1; PID:91370302 Aperimental source: strain 8288C

Generica: GGD NABLB; YGT2
Gene: SGD NABLB; YGT2
(Cross references: SGD:80004038; HIPS:YLR048v
HAp position: 12R
(Antrops: 3/0; Vesst ribosomal protein Sl.e

Query Match 1.0%; Score 7; DB 2; Best Local Similarity 100.0%; Pred. No. 58; Matches 7; Conservative 0; Mismatches 388 TEWAEEN 394 ||||||| | 240 TEWAEEN 246 0; Indels Length 252; 0 Gaps

0

RESULT 31
073236
hypothetical protein TM1079 - Thermotoga maritima (strain MSB8)
c;Species: Thermotoga maritima
c;Species: T Nature 399, 323-329, 1999 A;Title: Evidence for lateral gene transfer between Archaea and Bacteria from genome sequence of Thermotoga maritima

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pothetical protein N02B1.4 - Caenorhabditis elegans Species (aenorhabditis elegans Nate: 15-Oct.1999 sequence\_revision 15-Oct.1999 #text\_change 15-Oct.1999 Victosion: 713675 lightning, J.
maitted to the EMBL Data Library, October 1996
deference number: Z19780
cccssion: T23675

xry; translated from GB/EMBL/DDBJ

atus: preliminary: translated from GB/EMBL/DDBJ Jecule type: DM sidues: 1-313 AWIL-coms references: EMBL:281102; PIDN:CAB03203.1; GSPDB:GN00022; perimental source: clone NOZB1

CESP: M02B1.4

h;Hap position: 4 h;Introns: 31/2; 195/3; 228/1 CESP: M0281.4

Query Match 1.0%; Score 7; DB 2; Best Local Similarity 100.0%; Pred. No. 71; Matches 7; Conservative 0; Mismatches 0; Indels Length 313; 0 Gaps

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370 RTILFAS 376 ||||||| 91 RTILFAS 97

pcF10

a region of plasmid pCF10 containing positive control genes and

robable pheromone-responsive regulatory protein X - Enterococcus faecalis plasmid |
Species: Briefrococcus faecalis |
Species: Briefrococcus faecalis |
Species: Briefrococcus faecalis |
Species: Briefrococcus faecalis |
Species: Do-Jun-1991 |
Species: Do-Jun-1991 |
Species: Do-Jun-1991 |
Species: Do-Jun-1991 |
Species: Di-Jun-1991 |
Species: plasmid

Query Match 1.0%: Score 7; DB 2; Length 317. Best Local Similarity 100.0%; Pred. No. 71;

Page 41

NO.

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Page 42

A:Reference number: A72200; MUID:99287316 A:Accession: G72256 A:Status: Preliminary A:Motecule type: DNA A:Residues: 1-254 (ARM) A:Cross-references: GB:AE001768; GB:AE000512; NID:94981619; PIDN:AAD36155.1; PID:94981624; TIGR:TM1079 A:Cross-references: GB:AE001768; GB:AE000512; NID:94981619; PIDN:AAD36155.1; PID:94981624; TIGR:TM1079 C:Genetics:

Genetics: Gene: TM1079

ouery Match 10.0%; Score 7; DB 2; Length 264; Best Lock 60; Best Lock 60; Mismatches 0; Indels Matches 0; Indels 0

δÃ 516 ELVEKFY 522 ||||||| 16 ELVEKFY 22

RESULT 32 T24824

hypothetical protein 711A5.5 - Caenorhabditis elegans C:Species: Caenorhabditis elegans C:Date: 15-oct-1999 #sequence\_revision 15-oct-1999 #text\_change 15-oct-1999 C:Accession: 72.4824 R:Meditray A. Bull Data Library, May 1996 A:Reference number: 219939 A:Reference number: 219939

ttus: preliniary; translated from GB/EMBL/DDBJ legula type: DMA Lidnes: 1-286 MBL.272515; PIDN:CAA36685.1; GSPDB:GN00023: CESP:T11A5.5 yes\_references: BMBL.272515; PIDN:CAA36685.1; GSPDB:GN00023: CESP:T11A5.5 perimental source: close T11A5

A:Experimental source: C:Genetics: A:Gene: CESP:T11A5.5 A:Map position: 5 A:Introns: 202/2

Query Match 1.0%; Score 7; DB 2; Best Local Similarity 100.0%; Pred. No. 65; Matches 7; Conservative 0; Mismatches Length 288 0; Indels 0 Gaps

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₽ δÃ 442 EGFECKS 448 ||||||| 253 EGFECKS 259

RESULT

Apr 9 11:14:14 2001 Matches 7; Conservative 600 SKFSERL 606 |||||||| 48 SKFSERL 54 0 Mismatches us-08-403-803-2\_copy\_45\_750.olig.xpx 0 0 Gaps

Page 44

Diserved hypothetical protein - Deinococcus radiodurans (Strain R1)
Diserved hypothetical protein - Deinococcus radiodurans
Specias: Deinococcus radiocus radiodurans
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terence number: A75250; MID: 20036896

A:Status Preliminary
A:Molecule type: DAND
A:Residuse: 1518 (AND)
A:Residuse: Series: 0812
A:Residuse: 0812
A:Resi

overy Match 100.0%; Score 7; DB 2; Length 318; Best Local Similarity 100.0%; Pred. No. 72; Matches 7; Conservative 0; Mismatches 0; Indels

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Gaps

0

RESULT 36 T25869

structural gene

hypothetical protein 7094 / - Cannorhabdiis elegans Cipetcies: Cenorhabdiis elegans Cipetci 15-oct-1999 #sequence\_revision 15-oct-1999 #text\_change | C.Accession: 72869 | C.Acce 04-Mar-2000

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RESULT 40
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AlPolaculars 1.342 (AM)
AlCross-references: EMBLU97405; PIDN:AMB53009.1; GSPDB:GN00019; CESP:T09B4.7
AlExpertimental outce: strain Bristol N2; clone T09B4
C:Genetics: T08B4 7
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Introna: 58/1, 121/3; 226/1; 280/3
Superfamily: Kinase-related transforming protein; protein kinase homology
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284 TKKSPSP 290
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                                                                           Mon Apr 9 11:14:14 2001
RESULT 43
A71479
A71479
probable od shape protein - Chiamydia trachomatia (serotype D, strain UM3/CX)
C:Species: Chiamydia ttachomatia
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Db 16 SLFSAVK 22
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Phillips, C.A.; Richardson, D.; Heidelberg, J.; Sutton, G.G.; Plei
                                                                                                                                                                                                                                                                                                                                                                                                                                                      380 EEFGLLG 386
|||||||
282 EEFGLLG 288
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282 EEFGLLG 288
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cell division protein fiz2 C9056 [imported] - Campylobacter jejuni (strain NCTC 11168)
C.Space 131-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 19-May-2000
C.Noccession: C81340
R.Parkhill, J.; Wren, B.W.; Mungall, K.; Ketley, J.W.; Churcher, C.; Basham, D.; Chillingworth, T.; Davies, R.M.; Feltwell,
C.W.; Quall, W.; Rejandream, M.A.; Rutherford, K.M.; VenTilet, A.; Whitehead, S.; Barrell, B.G.
M.Tille: The genome sequence of the food-borne pathogen Campylobacter jejuni reveals hypervariable sequences.
A; Reference number: A81250; MUID: 20150912
A; Roccession: C81340
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A/Title: Evidence for lateral gene transfer between Archaea and Bacteria from genome sequence of Thermo A; Reference number: A72200; MUID:99287316

A; Rectance number: A72200; MUID:99287316

A; Status: Preliminary
A; Status: Preliminary
A; Status: Preliminary
A; Residues: 1-370 CARN
A; Residues: 1-370 CARN
A; Residues: 1-370 CARN
A; Cross: references: GB:AE001771; GB:AE000512; NID:94981678; PIDN:AAD36211.1; PID:94981682; TiGR:TM1135
A; Cross: references: GB:AE001771; GB:AE000512; NID:94981678; PIDN:AAD36211.1; PID:94981682; TiGR:TM1135
A; Cross: references: GB:AE000512; PIDN:AAD36211.1; PID:94981682; TiGR:TM1135
A; Cross: References: GB:AE000512; PIDN:AAD36211.1; PID:94981682; TiGR:TM1135
A; Cross: References: GB:AE000512; PIDN:AAD36211.1; PID:94981682; TiGR:TM1135
A; Cross: References: Referenc
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C;Superfamily: cell division protein fts2
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;Molsoula type; DNA
;Crossistes; 1-370 cPAR;
;Cros
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us-08-403-803-2_copy_45_750.olig.rps
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Page 48
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RESCUT. 42
H83/74.
H83/74.
H83/74.
H83/74.
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H83/74.
H83/74.
H83/74.
H83/74.
C.Species: Chlamydia muridarum, Chlamydia trachomatis McPn
C.Date: 31:Mar-2000 #sequence_revision 31:Mar-2000 ftext_change 02:Sep:2000
C.Accession: H83/74.
K.Read, T.D.; Brunham, R.C.; Shen, C.; Gill, S.R.; Heidelberg, J.F.; White, O.; Hickey, E.K.; Peterson, J.; Otterback, T.;
K.C.; Doddon, R.; Gvinn, M.; Nelson, W.; DeBoy, R.; Kolonay, J.; McClarty, G.; Salzberg, S.L.; Eisen, J.; Fraser, C.K.
K.Tilte: Genome sequences of Chlamydia trachomatis McPn and Chlamydia pneumoniae AR39.
A. Rectance number: A81500; MCID:20150355
A. Scottania (M.).
A. Scottani
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A.Experimental source: strain AR39, HL cells
C:Senetion:
A.Gene: CP1002
A.Gene: CP1002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C; Superfamily: rod shape-determining protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .;Status: preliminary
;Molecule type: DNA
;Molecule type: DNA
;Masiduse: 1-379 cTRD
;Mas
              Query Match 1.0%; Sometive 1.00.0%; In Matches 7; Conservative 0;
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Best Local Similarity 100.0%;
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              Score 7; DB 2;
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11:14:14 2001
                                                                                                                                                                                                                                                                                                                                       probable apore coat polysaccharide biosynthesis protein (apaC) - syphilis spirochete
c)species: Trepomena pallidum subsp. pallidum (syphilis spirochete
c)species: A:-Unity98 seed.com.
c)species: A:-Unity99 seed.com.
c)spec
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Ciphers D5-pecies Archaeoglobus tulgidus
Ciphers D5-pecies P5 (1997) sequence_revision 05-pec-1997 stext_change 22-Jun-1999
Cichers D6-pec-1997 sequence_revision 05-pec-1997 stext_change 22-Jun-1999
Ritlenk, H.P.; Clayton, R.A.; Tomb, J.F.; White, O.; Nelson, K.E.; Ketchun, K.A.; Dodson, R.J.; Gwinn, M.; Hickey, E.K.; Pec (1997) sequence_revision 05-pec-1997 stext_change (1997) sequence_revision 05-pec-1997 stext_change (1997) sequence_revision 05-pec-1997 stext_change (1997) sequence_revision (1997) sequence_revision 05-pec-1997 stext_change (1997) sequence_revision (1997) sequence_revi
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C; Superfamily: rod shape-determining protein
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A:Residues: 1-389 GCIM-
A:Cross:reterences: GS:AEO03992; GB:AEO03949; NID:g9105243; PIDN:AAF83217.1; GSPDB:GN00128; XFSC:XF0407
A:Cross:reterences: GS:AEO03992; GB:AEO03949; NID:g9105243; PIDN:AAF83217.1; GSPDB:GN00128; XFSC:XF0407
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N:Alternate names: hypothetical protein YRR0718 hypothetical protein YRR074v
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C:Species: 17-May-1994 sequence_revisiase
C:Accession: 84347; 845937; 859707; 837234
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R:yan der Aarty, Q.J.M: Barthe, C.; Doisnon, F.; Aigle, M.; Crouset, M.; Steensma, H.Y.
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[imported] - Xylella fastidiosa (strain 9a5c) percies: Xylella fastidiosa (species: Xylella fastidiosa consequence (species: Xylella fastidiosa consequence (species: 18-va-9000) asequence_revision 20-Aug-2000 stext_change 02-Sep-2000 (species: Xylella fastidiosa Consortium of the Organization for Nucleotide Sequencing and Analysis, Sao Paulo, Brasi (anonymous, The Xylella fastidiosa Consortium of the Organization for Nucleotide Sequencing and Analysis, Sao Paulo, Brasi (anonymous, The Xylella fastidiosa).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Title: The genome sequence of the plant pathogen Xylella fastidiosa. Reference number: A2515, MJID:20165717 Note: for a complete list of authors see reference number A59328 below Accession: F82809
                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        coss-references: PMBL.X76294, NID:9974203; PIDN:CAAS3931.1; PID:e264511; PID:9433938 recessing, NIC: you der Anri, Q.J.M. abmitted to the EMBL Data Library, November 1993 litted to the Protein Sequence Database, August 1994 (ference number: 9493)2 (geston: 8493)3 (gesto
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Query Match 1.0%; Score 7; DB 2; Length 415; Best Local Similarity 100.0%; Pred. No. 92; Matches 7; Conservative 0; Mismatches 0; Indels 0; Indels 0; Gaps 0

380 EEFGLLG 386 ||||||| 318 EEFGLLG 324

RESULT 50 G75363

integral membrane protein, NBAMP family - Deinococcus radiodurans (strain R1)
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Ribhite, 0.; Eisen, J.A.; Haidelberg, J.F.; Hickey, E.K.; Peterson, J.D.; Dodson, R.J.; Haft, D.H.; Gwinn, M.L.; Nelson, W.
M.; Shrn, M.; Vanathever, J.J.; Lam, P.; KdDonald, L.; Utterback, T.; Zalewski, C.; Makarova, K.S.; Aravind, L.; Daly, M.
S.; Smith, H.O.; Vonter, 1999.; Fraser, C.M.
Science 266, 1271-1577, 1999.; Fraser, C.M.
A:Pitle: Genome becquence of the radioresistant bacterium Deinococcus radiodurans R1.
A:Beterone muberquence of the radioresistant bacterium Deinococcus radiodurans R1.
A:Beterone muberquence (1975) MUD: 20036896

Page 53

A:Status; preliminary
A:Status; preliminary
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A:Residues: 1-456 (9817)
A:Cross-references GB-AE002012; GB:AE000513; NID:96459473; PIDN:AAP11265.1; PID:96459478; TIGR:DR1709; GSPDB:GN00077
A:Cross-references GB-AE002012; GB:AE000513; NID:96459473; PIDN:AAP11265.1; PID:96459478; TIGR:DR1709; GSPDB:GN00077
A:Cross-references GB-AE002012; GB-AE002013; NID:96459473; PIDN:AAP11265.1; PID:96459478; TIGR:DR1709; GSPDB:GN00077
A:CROSS-REFERENCES GB-AE002013; PID:96459478; TIGR:DR1709; GSPDB:GN00077

Ouery Match 2016 10%; Score 7; DB 2; Length 436; Best Local Similarity 100.0%; Pred. No. 96; Matches 7; Conservative 0; Mismatches 0; Indels 0; Indels 0;

Gaps 0

Ϋ́ 484 LGIASGR 490 ||||||| 95 LGIASGR 101

Search completed: April 6, 2001, 16:57:03 Job time: 151 sec

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Sequence 1, Appli 115	Mon Apr 9 11:14:12 2001 us-08-403-803-2_copy_45_750.clig.rai	7 19 2.7 19 2 .7 19 2 .8 cequence 1. Appl 9 17 2.4 22 19 305.08-394.152A-11 10 17 2.4 22 19 305.08-395.53-7 10 18 2.5 394.152A-7 10 18 2.5 394.152A-7 10 18 2.5 394.152A-7 10 18 2.5 394.152A-7 11 18 2.5 394.152A-7 12 39 1.3 32 394.152A-7 13 394.152A-7 14 39 1.3 395.394.152A-7 15 394.152A-7 16 394.152A-7 17 395.394.152A-7 18 394.152A-7 18 394.152A-7 19 395.394.152A-7 19 3	Mon Apr 9 11:14:12 2001 us-08-403-803-2_copy_45_750.olig.zal

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: LOCATION: 1.707
OTHER IMPORMATION: /mote* *EXTRACELLULAR DOMAIN OF
: OTHER IMPORMATION: PSMA*
PCT-US95-11720-18
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Best Local Similarity 100.0%; P
Matches 706; Conservative 0;
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Best Local Similarity 100.0%; Pred. No. 0;
Matches 706; Conservative 0; Mismatches
                                                                                          241 EANGLESIEVEID FORTDAOKLEKINGSSAPDISSERGSLKVEK MICHGEFORESTORVEK 300
245 EANGLESIEVEID FORTDAOKLEKINGSSAPDISSERGSLKVEKINGEGFORESTORVEK 344
301 HISTORYTEIT KNICTLEGAVERDEKVELGGHOUSENGSLKVEKINGEGFORESTORVEK 344
301 HISTORYTEIT KNICTLEGAVERDEKVELGGHOUSENGSLKVEKINGER STORVEKENGSTORVEK
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225 VILYSDPADYEAPCYKSYPDCHNILPGGGYORGNITLHJAGAGDPLTPGYFANETAYERGIA 284
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165 SPOGNPSCDLYVYNARTEDFFILERDMINGSCKIVLARYGYVFRONKYRAOLAGAKG 180
                                                                                                                                                                                                                                                                                                           100 LISVELARVOVLLSYPRKTHENKISIINEIXORIFWRISLFEPPPROXENVSDIVPPF8AF 120
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121 SPOGNEDSDLWYWNARTEDFFKLERDMKINGSGKTVTARYGKWEGGKWENDAGLAGAKG 180
122 SPOGNEDSDLWYWNARTEDFFKLERDMKINGSGKTVTARYGKWERGKWENDAGLAGAKG 181
                                                                                                                                                                                                                                                                                                                                                                           IOR APPLICATION DATA:
APPLICATION NUMBER: US/07/973,337A
APPLICATION NUMBER: US NOV 1992
FILING DATE: 05 NOV 1992
TORNEY/ACENT INFORMATION:
NAME: White, John P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      62 LOSVELAHYDVLLSYPNKTHPNYISIINEDGNEIFNTSLFEPPPPGYENVSDIVPPFSAF 121
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LOGY: linear
LE TYPE: protein
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Pred. No. 0;
; Mismatches 0; Indels
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US-08-135-53-2
US-08-135-53-2
Sequence 2. Application US/08325553
Patent bo. 5518866
Patent No. 5518866
REERAL IMPORATION:
APPLICANT: Heaton, Harten D. M.
APPLICANT: Heaton, Harten R.
APPLICANT: Fair, Miliam R.
APPLICANT: Fair, Miliam R.
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CHROTER NO.0. LE PORH:

MEDIUM TIE B. FLOPY disk
COMPUTENT E. B. FLOPY disk
COMPUTENT E. B. FLOPY disk
COMPUTENT E. B. FLOPY disk
COMPUTENT STEEL P. P. FLOSA FLOS
SOFTHALE STEEL P. P. FLOSA FLOS
SOFTHALE CONTROL IN Release $1.0, Version $1.25
COMBETY ADDITION HUMBER: MAD $1.25
FILING DARE: WA FEE 9.08/394,152A
FILING DARE: WA FEE 9.08/394,152A
FLOSEFICATION HUMBER: WAS 5.5
ATTORNEY/AGENT INFORMATION:
ANAME: White, John P.
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-08-194-152A-2
sequence 2, Application US/08394152A
satent No. 5935818
GENERAL INFORMATION:
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NPLICANT: Heaton, Murran D.W.
NPLICANT: Palt. William R.
NTTE OF INVESTION: PROSTRE-SPECIFIC MEMBRANE ANTIGEN AND
NTTE OF INVESTION: USES THEREOF
NUMBER OF SEQUENCES: 4
COMRESPONDENCE ADDRESS:
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525 FORLICIASGRARTKAMETNAFSGYPLYHSVYETYELVEKFYDPMFKYHLTVAQVRGGNY 594
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JIP: New York
STATE: New York
THEE: New York
THEE: New YORK
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362 LKEGHAPBRILIHANHULAEERGLLGSTEMAEENSKLLØERGYAYINADSSIEGNTILEN
363 LKEGHAPBRILIHANHULAEERGLLGSTEMAEENSKLLØERGYAYINADSSIEGNTILEN
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BDIUM TYPE: Floopy disk

DENIM TYPE: Floopy disk

PERMING SESSEM: DC-DOS/MS-DOS

OFTWARE: PATENTING REMESS #1.0, Version #1.25

REWIT APPLICATION NUMBER: US/08/325,553
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EET: 1185 Avenue of the Americas
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RESSEE: Cooper & Dunham
EET: 30 Rockefeller Plaza
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TY: Balt, William R.

TIVENITOM: THE PROSTATE-SPECIFIC MEMBRANE ANTIGEN

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REFERENCE/DOCKET NUMBER: 41426-B
REFERENCE/DOCKET NUMBER: 41426-B
REFERENCE/DOCKET NUMBER: 20.00
TRIEPRAN R: 2123 37.00-00
TRIEPRAN R: 2123 391-0525
INVOHATION ROY SEQ ID No. 2:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 100.0%; P
Matches 706; Conservative 0;
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TELECOMPUNE_ACTION INFORMATION
TELEPHONE: 650-499-595
TELEPAX: 66141 PENNIE
TELEPAX: 66141 PENNIE
TELEPAX: 66141 PENNIE
TELEPAX: 750-4810-6148
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215 VILYSDPADPRAPGVRSYPDGNILPGGGVORGNILNLNGAGDPLPFGYPANEYAYRSGIA 284
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165 SPOGNESSDLYFYNYAFESPFELESBMEINSGELYTAR GRYPGGRYCHAGAGAGG 224
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465 DCTPLAYSLVINLTKELKSPDEGEGKSLYESMTKKSPEPEFSCAPFRISKLGGODERVF 524
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405 LKECHEPRETILPASHDAEERGLLGSTEWAEERSLLGEGVAX INADSSIEGUNTLEW 464
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285 EAVGLPSIPHIPTOYYDAQKLLEKMGCSAPDSSHROSLKYPYMGPGPTGNESTQKYKX 344
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285 EAVGLESIPHEPGYTDAQKLLERMGGSAPPDSSHRGSLKYPYRVGPGFTGNESTQKVIM 344
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165 SPOGMPEGDLYPVNYARTEDFFKLERDMKINGSGKIVLANGKVFRGKKVENDLAGAKG 224
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                421 DCTPLAYSLVINLY,RELKSPDEGEFECKSLYESHTIKESPSPESGAPRISKEGSGADFEVF 480
465 DCTPLAYSLVINLYRELKSPDEGEFECKSLYESHTIKESPSPESGAPRISKEGSGADFEVF 524
                                                                                   361 LAKEGHRPHRTILPASHDAEPGLLGSTEMAEENSRLLGERGVAY INADSSIEGNYTLRV 420
405 LAKEGHRPHRTILPASHDAEPGLLGSTEMAEENSRLLGERGVAY INADSSIEGNYTLRV 464
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% TYPE: No. 6150508e
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TYPE: protein
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Pred. No. 0;
); Mismatches 0; Indels
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ETOUTHEY DEAL FORM.

ETOUTHEY REPORT FORM.

COMPTER REPORT DIASETTE.

ETITAL DATE: 16-MAR-1996

PRIOR APPLICATION HOMBER:

ETITAL DATE: 16-MAR-1996

PRIOR APPLICATION HOMBER:

ETITAL DATE: 16-MAR-1996

PRIOR APPLICATION HOMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 11, Application US/08325553
Patent No. 5538866
GENERAL INFORMATION:
                                                                                                                          APPLICANT: Israeli, Ron S.
APPLICANT: Fair, William R.
TITLE OF INVENTION: THE PROSTATE-SPECIFIC MEMBRANE ANTICEN
NUMBER OF SEQUENCES: 38
CORRESSEE: Cooper's Dunham
STREET: 30 Rockefeller Plaza
CITY: New York
COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-325-553-11
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APPLICANT, BOLTO, ALTON D.

APPLICANT, HOLTO, ETCH H.

APPLICANT, HOLTO, ETCH H.

APPLICANT, HOLTO, BLILLAN THOMAS

APPLICANT, THE MERCHANT AND PROSTATE-SPECIFIC TITLE OF INVESTION. FOR THE EXTRACELUSIAN DOMAIN OF PROSTATE-SPECIFIC NUMBER OF INVESTION. MEMBRANE ANTIGEN

ADDRESSED OF ROPERIORS J.

ADDRESSED PROMISES Edmonds, LLP

STREET; 1155 Venue of the Americas
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481 PORICIASCRARYTRINEPINKESCYPLYBSYETYELVEKEYDPHEKTHIJVAVGGON
525 PORICIASCRARYTRINEFINKESCYPLYBSVETYELVEKEYDPHEKTHIJVAQVBGGON
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705 FPGIYDALFDIESKVDPSKAMGEVKRQIYVAAFTVQAAAETISEVA 750
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601 KESERLODEDKSHETVLROMNOLAFIZBARTIDELGIJDEDEFFRHYTYASSSHRYKAGES 660
601 KESERLODEDKSHETVLROMNOLAFIZBARTIDELGIJDEDFFRHYTYAPSSHNRYAGES 704
CLASSIFICATION: 435
PRICH APPLICATION DATA:
APPLICATION NUMBER: U9/07/973,337A
PILHE OMET INFORMATION:
AMET: WANTE, JOHN P.
BRISTANTON WHOBER: 29,678
BRISTANTON WHOBER: 1747/41426
REFERENCE/DOCKET NUMBER: 1747/41426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          601 KESEKLOPEDKSHPYULHMINDOLMFLERNFIDPLGLEDREPTHYTAPSSHHKYAGES
645 KESEKLOPEDKSHPYULHMINDOLMFLERNFIDPLGLEDREPTHYTAPSSHHKYAGES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           541 FELANSIVLPFDCRDYAVVLRKYADKIYSISMKHPQENKTYSVSFDSLFSAVKNFTEIAS 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
TORNEY/AGENT INFORMATION:
NAME: Baldwin, Geraldine F
REGISTRATION NUMBER: 31,232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New York
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TELEPHONE: (212) 977-9550
TELEPKA: (212) 64-0525
TELEX: 42523 COOP UI
MANTON FOR ESD ID NO. 11:
UNMATION FOR ESD ID NO. 12:
LENGTH: 19 main acids
TYPE: anino acid
TYPE: double
ODOLOGY: linear
DOULOGY: linea

ORGANISM: Homo sapien
TISSUE TYPE: Carcinoma
IMMEDIATE SOURCE:
CLOME: Prostate Specific Membrane Antigen
US-08-325-553-11 SOURCE

Ouery Match 2.7%; Score 19; DB 1; Length 19; Best Local Similarity 100.0%; Pred No. 2 &et 2; Matches 19; Conservative 0; Mismatches 0; Indels

Gaps

CLONE: Prostate Specific Membrane Antigen US-08-394-152A-11

RGANISM: Homo sapien ISSUE TYPE: Carcinoma EDIATE SOURCE: RGANISM: "

Query Match 2.7%: Score 19: DB 2: Length 19: Best Local Similarity 100.0%: Pred. Mo. 2.6e-1; Matches 19: Conservative 0: Mismatches 0; Indels

0 Gaps

0

656 YAGESFPGIYDALFDIESK 674

-08-622-0468-21 Sequence 21, App Patent No. 6103 GENERAL INFORM

Application US/08622046B

LICANT: Saedi, Mohammed S. LICANT: Mikolajozyk, Stephen D. ILBO OF INVENTION: Stable Variant HK2 Polypeptide IBER OF SEQUENCES: 21 RESPONDENCE ADDRESS:

Schwegman, Lundberg, Woessner & Kluth, P.A O. Box 2938

1, Application US/08394152A 5935818

WT: Heston, Warren D.W.

"Y: Fair, William R.
INVENTION: DESCRIPTION DEMORANE ANTIGEN AND LIVENTION: USES THEREOF

P SOUTHCES: 48 THEREOF

PP SOUTHCES: 48 THEREOF

United States of America Cooper & Dunham LLP 5 Avenue of the Americas

MEDIUM TYPE: Floppy disk COMPUTER: IBM 330 466 DX2 OPERATING SYSTEM: PC-DOS/MS-DOS

Mon Apr 9 11:14:12 2001

MBER: US/08/622,046B 26-MARCH-1996

TREATMONE (10.14-3)7-5940

INFORMATION FOR EDRY IN NO. 2
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
THE WITH: 10 anino acids
STREATMONESS: single
TORACCE TIPE: Peptide
US-08-627-0468-21

ANT: HESSON, WAITEN D.W.
ANT: HESSON, WAITEN D.W.
ANT: HESSON, WAITEN D.W.
ANT: BAIN WILLIAM R.
OF INVENTION: THE PROSTATE-SPECIFIC MEMBRANE ANTIGEN
OF SEQUENCES: 38
DESCREE: Cooper & Dunham
ESSEE: Cooper & Dunham
ESSEE: Cooper & Dunham Application US/08325553 5538866

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Floppy disk
M PC compatible
M PC pOS/MS-DOS
EentIn Release #1.0, Version #1.25
IION DATA:

: Woessner, Warren D.
STRATION NUMBER: 30,440
RENCE/DOCKET NUMBER: 476.001US1
MMUNICATION INPORMATION:
MFUNICATION: 612-373-6503

152 CSGKIVIARYGKVFRGNK 169

7-3061 ID NO: 21:

Query March 2.5%; Score 18; DB 3; Length 18; Best Local Similarity 100.0%; Pred, Mo. 2.6e-1; Onservative 0; Mismarches 0; Indels

Page 31

Mon

Apr 9 11:14:12 2001

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Page 32

CITY: New York
STATE: New York
COUNTRY: United States of America
IP: 10112
IP: 1012

ICATION: 435
LICATION DATA:
TION NUMBER: US/07/973,337A
DATE: 05 NOV 1992
ACENT INFORMATION:
White, John P. 1747/41426

JATE:
//ACEPY INFORMAT:
//ACEPY INFORMAT:
//ACEPY INFORMAT:
MILTE, JOHN P.
MILTE, JOHN P.
MILTE, JOHN P.
MILTE, JOHN MUNEER: 128.678
AERNCE/COCKET NUMBER: 128.7474.1
AMENICATION HUNBER: 128.755
ALERNA: (121) 664-0553
ALERNA: (121) 674-053
ALERNA: (1

ORICIPAL SOURCE:
ORGANISM: Homo sapien
TISSUE TYPE: Carcinoma
IMMEDIATE SOURCE:
LOUNE: Prostate Specific Membrane Antigen
US-00-125-553-7

Query Match 2.4%; So Best Local Similarity 100.0%; I Matches 17; Conservative 0; Score 17; DB; Pred. No. 3.2 0; Mismatches DB 1; 140. 40. 3.2e-10; 0;

Length 22; Indels

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Ŷ 34 TQIPHLAGTEQNFQLAK 50

RESULT 10

Apr 9 11:14:12 2001

ARE: Patentin Release \$1.0, Version \$1.25 APPLICATION DATA: ATION NUMBER: US/08/394,152A

24-FEB-95 NFORMATION:

PRICE/CONTENT NUMBER: 28,678
RUNICATION INFORMATION:
HUNICATION INFORMATION:
HORE: (212) 29-0400
X: (212) 39-0523
Y FOR SEQ ID NO: 11:
CHARACTERISTICS:
19 amino.

mino acid DNESS: double : linear

NO NO

Page 29

Mon

us-08-403-803-2\_copy\_45\_750.olig.rai

wence 7, Application US/08394152A int No. 5935818

ISTANELI, ROD S.
Heston, Warren D.W.
FRAIT, WILLIAM R.
FRAIT, WILL

H TYPE: FI

:: Ploppy disk
IBM 330 466 DX2
IBM 330 466 DX2
POSTEM: PC-DOS/MS-DOS
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CATTON DATM:
(NUMBER: US/08/394,152A

-νποΕR: US/08/394,152A 24-FEB-95 1: 436

United States of America

Cooper & Dunham LLP Avenue of the Americas

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Page 34

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TISSUE TYPE: Carcinoma
IMMEDIATE SOURCE:
CLONE: Prostate Specific Membrane Antigen
US-08-394-15JA-9
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ZIP: 10112

OUPTIER ROLLS

COMPUTER STATE

COMPUTER: 150 PC compatible

COMPUTER: 150 PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTMARE: Patentin Release #1.0, Version #1.25

""" APPLICATION DATA:

""" APPLICATION DATA:
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US-08-32-553-9
Sequence 9, Application US/08325553
: Patent No. 5538866
: Patent No. 55388670N:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Apr 9 11:14:12 2001
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Best Local Similarity 100.0%; P
Matches 16; Conservative 0;
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AX: (213) 391-0525
ION FOR SED ID NO: 9:
CE CHARACTERISTICS: 37H: 17 amino acids
E: amino acids
E: amino acid
ANNUEDNESS: double
POLOCY: linear
COULE TYPE: peptide
OTHETICAL: NO
OTHETICAL: NO
II-SENSE: NO
II-SENSE: NO
II-SENSE: NO
III-SENSE: NO
II
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TISTERNICO NUMBER: 174/41426

TOMMUNICATION INFORMATION:
LEPRAN: (212) 664-0525

LEX. (212) 664-0525

LEX. 42523 COOP UI

VITON POR SEQ ID NO: 9:
SNCE CHARACTERISTICS:
SNCE CHARACTERISTICS:
SNCE CHARACTERISTICS:
SNCE CHARACTERISTICS:
SNCE CHARACTERISTICS:
TER READABLE FORM:
TER READABLE FORM;
BUTTER: 12H PC compatible
SEATING SYSTEM: PC-DOS/MS-DOS
FTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                           COANT: HERED, MATTEN D.W.
COANT: HERED, MATTEN D.W.
COMPIT PAIT, WILLIAM R.
COF INVESTICE: THE PROSTATE-SPECIFIC MEMBRANE ANTIGEN
SEPONDENCES: 38
SEPONDENCE ALDRESS: ...
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IMPLICATION DATA:
ICATION NUMBER: US/07/973,337A
IG DATE: 05 NOV 1992
IX/AGENT INFORMATION:
IX/AGENT INFORMATION:
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New York
RY: United States of America
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30 Rockefeller Plaza
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38866
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EMTION: THE PROSTATE-SPECIFIC MEMBRANE ANTIGEN
QUENCES: 38
                                                                                                                                                                                                                         Inited States of America
                                                                                                                                                                                                                                                                                                                              Cooper & Dunham
Rockefeller Plaza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        eston, Warren D.W.
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Pred. No. 2.6e-09;
O; Mismatches O; Indels
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Page 36

Mon Apr

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Page 35

Query Match

2.44;

Score 17;

DB 2; Length 22;

Prostate Specific Membrane Antigen

ANDEMATION.

A NUMBER: 28.678

A NUMBER: 41426-B

CATION INFORMATION

E. (212) 276-0400

MORE STREET

B. (212) 276-0400

MORE STREET

B. (212) 276-0400

MORE STREET

B. (212) 276-0400

MORE STREET

CHARCEBERISTIES

TH. 22 MAIN CALL

TRANSPERSES: double

"PORY: Inform

: CLONE: Prostate Specific Membrane Antigen US-08-325-553-9

ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Homo sapien
TISSUE TYPE: Carcinoma
IMMEDIATE SOURCE:

OLECULE TYPE: peptide

Query Match 2 3%; Score 16; DB 1; Length 17; Best Local Similarity 100.0%; Pred. No. 2.6e-09; Matches 16; Conservative 0; Hismatches 0; Indels

0, Gaps 0

181 VILYSDPADYFAPGVK 196

RESULT 12 US-08-394-152A-9

Application US/08394152A 5935818

. ISRAELI, RON S.
T. HERCON, WALTEN D. N.
T. WESTLON, LINCOLD N.
T. WESTLON, LINCOLD PROCEDIC MEMBRANE ANTIGEN AND
INTENTION. PROCEDITOR MEMBRANE ANTIGEN AND
SEQUENCES. GESS THEREOP
SEQUENCES. GESS. THEREOP
TO SECUENCE TO SECUENCE THE SECUENCE TO SECUE

EDTIN 1002 EDTIN 17PE: Ploppy diak EDTIN 17PE: Ploppy diak EDTIN 17PE: Ploppy diak EDTIN 17PE: PC-DOS/MS-DOS POPMARE: PARENTIA RELEASE #1.0, Version #1.25 SEET APPLICATION NATA: REPTIANCE NATE: 24-FEB-55 CLASSIFICATION: 435 CLASSIFICATION: 435 CLASSIFICATION: 435

NUMBER: 28,678 KET NUMBER: 41426-B

Y: United States of America 10036

NCE ADDRESS:
:: Cooper & Dunham LLP
:: 1185 Avenue of the Americas

APPLICATION DATA: CATION NUMBER: US/08/325,553

NUMBER OF SEQUENCES: 48
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper 6 Dunham LLP
STREET: 1185 Avenue of the Americas
CTTY: New York

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Mon Apr 9 11:14:12 2001
HYPOTHE...
AFT TRANSE: No
ORITE TRANSE:
ORGANIZATION SAPLEN
TISSUE TYPE: COUNCE:
IMMEDIATE SOURCE: Specific Membrane Antigen
CLOME: Prostate Specific Membrane Antigen
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US-08-394-152A-10
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TISSUE TYPE: Carcinoma
IMMEDIATE SOURCE: CARCINOMA
CLORE: Prostate Specific Membrane Antigen
US-08-325-553-10
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REFERENCE JOGO NUMBER: 7.26.678
REPERENCE JOGO NUMBER: 1047/41426
RELECOMUNICATION I WOMENTON:
TELEPHONE: (1.21.26.879.25.5)
TELER: (2.12.16.879.25.5)
TELER: (2.12.16.879.25.5)
TELER: (4.12.23.1000 U.S.
TELER: (4.12.23.1
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APPIJCANT: HEROON, WAITEN D. H.
APPIJCANT: PEAIT, MILIEM R.
TITLE OF INVENTION: PROSTATE-SPECIFIC MEMBRANE ANTIGEN AND
TITLE OF INVENTION: USES THEREDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ce 10, Application US/08394152A: No. 5935818
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OMMUNICATION INFORMATION I
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ANDEDNESS: double
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Watten D.W.
THE MATTER D.W.
THE MET MATTER PROSTATE-SPECIFIC MEMBRANE ANTIGEN
PROPRESS:
BENE CON---
TEE: CON---
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TYPE: Floppy disk
ER: INS PC compatible
ING SYSTEM: PC-DOS/MS-DOS
ING SYSTEM: PC-DOS/MS-DOS
APILICATION NATA-ass #1.0, Version #1.25
APILICATION NATA-ASS
ANTION NUMBER: US/08/325,553
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30 Rockefeller Plaza
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CAATION DATA:
CON NUMBER: US/07/973,337A
TE: 05 NOV 1992
EENT INFORMATION:
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United States of America
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No
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PEGANISH PUBL CATCINGMA
LIMITSHE SOURCE
CLARKES SOURCE
194-152A-10
US-08-394-152A-10
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Best Local Similarity 100.0%; Pred, Mo. 2.5e-08;
Matches 15; Conservative 0; Mismatches 0; Indels
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| AFIDPLSLPDRPFYR 15
                                                                                                                                                                                                                                                                              GE: White, John P.
ISTRATION NUMBER: 28,678
ISTRATION NUMBER: 41426-B
DOMUNICATION INFORMATION:
EPRONE: (212) 278-9409
EPRA: (212) 391-0525
EPRA: 5864 ED NO: 8:
WIGN FOR 5864 ED NO: 8:
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DIUM TUPE: E10ppy disk
NOUTER: LEM 330 466 DX2
EBATING SYSTEM: PC-DOS/MS-DOS
ETHARE: PALENTIN RELEASE $1.0, Version $1.25
BWT ARPLICATION BUTNAN
PLICATION NUMBER: US/08/394,152A
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IDM TITE: Floopy disk
PUTER: 15M 304 466 DX2
PUTER: 15M 304 466 DX2
RATING SYSTEM: PC-DOS/MS-DOS
TWARE: Patentin Release #1.0, Version #1.25
PT APPLICATION DATA:
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ANT: HESTON, MATTEN D.W.
ANT: FALT, WILLIAM R.
OF INVENTION: USES THERBOF
OF INVENTION: USES THERBOF
OF SEQUENCES: 48
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NCE/DOCKET NUMBER: 41426-B

NCE/DOCKET NUMBER: 41426-B

NDE: (12) 279-040

NE: (212) 391-0525

N: (212) 391-0525

N: (212) 391-0525

N: FOR SEQ ID NO: 10: CHARACTERISTICS: 15 aniho acids

amino acids
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1185 Avenue of the Americas
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E: peptide
NO
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FEB-95
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TISSUE TYPE: Carcinoma
IMMEDIATE SOURCE:
CLONE: CLONE:
CLONE: SOURCE:
US-08-394-152A-6
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ORICINAL SOURCE:
ORGANISM, Homo Sapien
CELL Type: Carcinoma
IMMEDIANE SOURCE;
IMMEDIANE SOURCE;
CLORE; Prostate Specific Membrane Antigen
US-08-394-152A-34
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Best Local Similarity 100.0%; Pred, No. 2,3-05;
Matches 12; Conservative 0; Mismatches 0; Indels
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1 GVILYSDPADYFAP 14
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1 RTILFASWDAEE 12
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int No. 553886

irral INFORMATION:

IPPLICANT: IMPRANT
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LE TYPE: peptide
ETICAL: NO
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PAX: (212) 391-0525
TON FOR SEG ID NO: 34:
CE CHARACTERISTICS:
CE CHARACTERISTICS:
CHARACTERISTICS:
CHARACTERISTICS:
CHARACTERISTICS:
CHARACTERISTICS:
CB CHARACTERIS
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APPLICATION DATA:
APPLICATION BOTA:
APPLICATION NUMBER: U5/07/973,337A
NG DATE: 05 NOV 1992
BY/AGENT IMPORMATION:
White, John P.
NMT: JETACLÍ, ROD S.
NMT: REACON, FASTRON D.W.
NMT: RELT, HILLED R.
PF INVENTION: THE PROSTATE-SPECIFIC MEMBRANE ANTIGEN
OF SEQUENCES: 39
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TYPE: Plopy disk
NG SYSTEM: PC-005/MS-005
NG SYSTEM: PC-005/MS-005
PETCOM Release $1.0, Version $1.25
PETCOM AUTON DATA PARTON PLANTS. US/08/325,553
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Reaton, Marten D.M.
Reat, William R.
NYEARTON: THE PROSTATE-SPECIFIC MEMBRANE ANTIGEN
SEQUENCES: 39
SERCE ADDRESS:
ESD GOOPET & Dunham
ESD GOOPET & Dunham
ESD GOOPET & STATE PLANE
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NUMBER: 41426-B
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CELL TYPE: Carcinoma
: IMMEDIATE SOURCE:
: CLANE: Prostate Specific Membrane Antigen
US-08-325-553-34
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                                                                                                                                                                                                                                                                                                                                                                            ANTI-SENSE: NO
ORIGINAL SOURCE:
ORANISM: Homo Sapien
TISSUE TYPE: Carcinoma
IMMEDIATE SOURCE:
CLORE: Prostate Membrane Specific Antigen
US-08-135-553-31
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Best Local Similarity 100.0%; F
Matches 12; Conservative 0;
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Best Local Similarity 100.0%; Pred. No. 1.3e+05;
Matches 9; Conservative 0; Mismatches 0; Indels
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REFERENCE_DOCKET NUMBER: 1747/41426
RELECOMONICATION INFORMATION
TELECHONE: (211) 977-9509
TELECHONE: (211) 964-0525
TELEX: (422) COOP UI
TELEX: (422) COOP UI
TELEX: (422) COOP UI
TELEX: (422) COOP UI
TENDRATICS INFORMATICS II
SEQUENCE CHANCTERISTICS II
LENGTH: 12 anino acids
TYPE: anino acid
STANDEDNESS: double
STANDEDNESS: double
TODOLOGY: Linear
MOLECULE TYPE: EPPtide
HYG/METICAL: NO
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AGENT INFORMATION P. 28.678

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MEDIUM TYPE: Floppy
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ADDRESSEE: Cooper & Dunham
RRESTREET: 30 Rockefeller Plaza
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APPLICATION DATA:
ICATION NUMBER: US/07/973,337A
NO DATE: 05 NOV 199
EY/AGENT INFORMATION:
ENTITE, John P.
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TER: IBM PC compatible.

TRIS SYSTEM: PC-DOS/MS-DOS

TRIS SYSTEM: PC-DOS/MS-DOS

TRIS PAtentin Release #1.0, Version #1.25

APPLICATION DATA: US/08/325,553

ATION NOMBER: US/08/325,553
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12: Cooper & Dunham LLP

1185 Avenue of the Americas

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FAIX, William R.
'SWYION: PROSTATE-SPECIFIC MEMBRANE ANTIGEN AND
TENTION: USES THEREOF
JUNNESS: 48
'E ADDRESS: 48
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Pred. No. 2.3e-05;
O; Mismatches O; Indels
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ouery Match 13%; Score 9; DB 2; Length 9; Best Local Similarity 100.0%; Pred. No. 1.3e+05; Matches 9; Conservative 0; Mamatches 0; Indel:

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Gaps

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Best Local Similarity 100.0%, Pred, No. 0.87;
Matches 9; Conservative 0; Mismatches 0; Indels
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CURRENT FILMS DATE: 1999-06-11
EARLIER FILMS DATE: 1999-06-12
EARLIER FILMS DATE: 1996-06-12
NUMBER OF SEQ ID NOS: 14
SOFTWARE: PARENTIN VET: 2.0
SED ID NO 1
LENGTH: 481
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-09-330-095-1
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9-044-668-1
9-044-668-1
tent No. 6150508
ENERAL INFORMATION:
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| 176 AEEFGLLGS 284
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TISSUE TYPE: Carcinoma
IMAEDIATE SOURCE:
CLONE: Prostate Membrane Specific Antigen
                   MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Mindows
SOPTWARE: FASTEM: Mindows
SOPTWARE: FASTEM: Mindows
Version 2.0b
MREBUT APPLICATION DATA:
MPDICATION NOBER: US/09/044,668
FILING DATE: 18-MAR-1998
CLASSITICHTION: 536
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PULCANT: Hatch I, William R.

PULCANT: HATCH I WILLIAM R.

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POPULCANT R.

POPULCANT
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YPE: amino acid
'TRANDEDNESS: double
'OPOLOGY: linear
LECULE TYPE: peptide
OTHETICAL: NO
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White, John P.
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ITPE: Floppy disk
IER: IBM 310 466 DX2
IER: IBM 310 466 DX2
IFFE: FROM 1000/MS-DOS
REPLICATION DATA: TO Version $1.25
NATION STATEM IN CONTROL ICA
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NTE: 24-FEB-95

NATION: 435
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United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  olmes, Eric H.
Thomas
T
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5 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     us-08-403-803-2_copy_45_750.olig.rai
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear TOPOLOGY: linear MOLECULE TYPE: No. 6150508e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 22
US-09-130-095-1
Sequence 1. Application US/09330095
Sequence 1. Application US/09330095
Patent No. 6127161
Patent No. 6127161
APPLICANT: Rickeman Comporation
TITLE OF INVENTION: Levelne Aminopeptidase Gene, Recombinant DNA,
TITLE OF INVENTION: Process for Producing Leucine Aminopeptidase
FILE REPERRNCE: Pr-022
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 21
US-09-330-055-6
US-09-330-055-6
Sequence 6, Application US/09330095
Patent No. 6172161
GENERAL INFORMATION:
APPLICANT: Kikkoman Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 24
US-08-1325-53-6
US-08-1325-53-6
Sequence 6, Application US/08325553
Patent No. 5538866
GENERAL INFORMATION:
GENERAL INFORMATION:
STREEL ROOMS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; ORGANISM: Aspergillus sojae
US-09-330-095-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 1.1%; Score 8; DB 3; Length 8; Best Local Similarity 100.0%; Pred. No. 1 3+05; Matches 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 maino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PERICANT: Kikkoman Corporation
TITLE OF INVENTIONS: Joudine Aminopeptidase Gene, Recombinant DNA,
TITLE OF INVENTIONS: Process for Producing Leucine Aminopeptidase
TILE REPERBENCE: PH-622
URRENT APPLICATION NUMBER: US/09/30,095
URRENT FILING DATE: 1999-06-11
MARLIER APPLICATION NUMBER: UF-164611/1998
MARLIER APPLICATION NUMBER: UF-164611/1998
MARLIER FILING DATE: 1999-06-12
UMBER: OF SEQ 1D NOS: 14
OFFMARE: PARCHIN Ver. 2.0
COMPUTER READABLE FORM:
MEDIDM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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10 AEEFGLLGS 18
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||||||||
1 TEONFOLAK 9
                                                                                                                                                                                                                                                                                                                                                                                                                               DICCANT: Israeli Ron S.
PLICANT: Heston, Warren D.W.
PLICANT: Fair, William R.W.
TILE OF INVENTION: THE PROSTATE-SPECIFIC
MEER OF SEQUENCES: 38
RRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
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TONNEY_AGENT_INFORMATION:

AME: Baldwin, Geraldine F

BEGISTANION NUMBER: 11,732

BETERBNCE_POOCKET NUMBER: 6511-0013-999

BECOMMUNICATION INFORMATION:

BECOMMUNICATION 1809RAMION:

BECOMMUNICATION 1809RAMION:
                                                                                                                                                                                                                                                                                                                                            : Cooper & Dunham
30 Rockefeller Plaza
                                                                                                                                                                                                                   United States of America
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PORCHINAL SOURCE: SOURCE: SOURCE: NO
PORCHINAL SOURCE: Arcinoma
IMPERIOR SOURCE: Arcinoma
CLANE: PORCHICE: Specific Membrane Antigen
US-08-325-553-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 1.1%; Score 8; DB 1; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.3e+65;
Matches 8; Conservative 0; Mismatches 0; Indels
             Owery Match 9: Conservative 0; Mismatches 0; Indels Matches 8; Conservative 0; Mismatches 0; Indels
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1 IYNVIGTL 8
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TETICAL: NO
SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CISTRATION NUMBER: 28,678
PERRENCE/DOCKET NUMBER: 114/41426
PERRENCE/DOCKET NUMBER: 1214/41426
COMMUNICATION INFORMATION
LEPHONE: (212) 977-9550
LEPHONE: (212) 646-0525
LEPA: 41253 COOP UI
LEX: 41253 COOP UI
VITON POR SED ID NO: 6:
NUCL CHARACTERISTICS:
NUCL CHARACTERISTICS:
                                                                                                                                                                                        X: 2647
XION FOR SEQ ID NO: 2:
MCE CHARACTERISTICS:
MCTH: 1577 amino acids
PE: amino acid
                                                                                                                   LOGY: not relevant
LE TYPE: protein
AL SOURCE:
NISM: Streptococcus salivarius
                                                                                                                                                                                                                                                                                              FICATION: 800
PPLICATION DATA:
ATION NUMBER: AU PM7643
ATION NUMBER: AU PM7643
S.DATE: 24-AUG-1994
GUNICATION IMPORMATION:
GUNICATION IMPORMATION:
GORE: 61 2 9957 5944
AX: 61 2 957 6288
                                                                                                                                                                                                                                                                                                                                                                                                                                     READALE PORM:
TITE: Elbpy disk
TER: IMM PC compatible
TWO SYSTEM: PC-DOS/MS-DOS
NPLICATION DATA
NATION NUMBER: US/08/793,824
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/08793824
5981838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TY: Israeli, Ron S.
TY: Heston, Warren D.W.
TY: Pair, William R.
INVENTION: PROSTATE-SPECIFIC MEMBRANE ANTIGEN AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RE: Patentin Release #1.0, Version #1.25
IPPLICATION DATA:
TION NUMBER: US/08/325,553
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35818
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Level 8, 168 Walker Street
o. 5981838th Sydney
New South Wales
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E: 05 NOV 1992
NT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ), Christine Lynn
3, Philip Morrison
5, Micholas Anthony
6 Genetic Manipulation of Plants to
Increase Stored Carbohydrates
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ORGANISH: Homo sapien
TISSUE TYPE: Carcinoma
INMEDIATE SOURCE:
CLOWE: Prostate Specific Membrane Antigen
US-08-194-1127-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 113: Score 8; DB 2; Length 9:
Best Local Similarity 100.0%; Pred. No. 1.3e+65;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SULT 4,
-08-325-553-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1091 AFLDELKA 1098
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 311 IYNVIGTL 318
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1 IYNVIGTL 8
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM 330 466 DX2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: USES THEREOF UMBER OF SEQUENCES: 48
                                                                                                                                                                                                                                                                                                                                                                                   RATING SYSTEM: PC-DOS/MS-DOS
TWARE: PatentIn Release #1.0, Version #1.25
NT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ANT. Heston, Walten D.M.

DET INVENTION HILLEM THE PROSTATE-SPECIFIC MEMBRANE ANTIGEN
OF SEQUENCES.
OF SEQUENCES.
DONORMUE ADDRESS:
SESSEE: GOODER, DUNDANE
SESSEE: GOODER, DUNDANE
() NAV YORK FEILER Place
                                                                                                                                                                                                                                                                                                   SIFICATION: 435
RAPLICATION DATA:
ICATION UNMER: US/07/973,337A
RG DATE: 05.NOV 1992
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New York
Y: United States of America
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TION NUMBER: 28,678
E./DOCKET NUMBER: 41426-B
E./DOCKET NUMBER: 41426-B
E: (212) 278-0400
E: (212) 391-0525
E/SCR SEQ ID NO: 6:
FOR SEQ ID NO: 6:
HARACTERISTICS:
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                                                                                                                                                                                           GENT INFORMATION:
hite, John P.
TION NUMBER: 28,678
TION NUMBER: 174
ECPOCKET NUMBER: 174
ICATION INFORMATION:
E: (212) 977-9550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  # York
United States of America
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24-FEB-95
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RESULT 28
US-08-394-152A-5
US-08-394-152A-5
I Sequence 5, Application US/08394152A
Patent No. 5035818
GENERAL INFORMATION:
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; TISSUE TYPE: Carcinoma
; IMMEDIATE SOURCE:
; CLONE: Protate Specific Membrane Antigen
US-08-32-531-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Owery Match 10.0%; Score 7; DB 1; Length 7; Best Local Similarity 100.0%; Pred. No. 1.3e+05; Matches 7; Conservative 0; Mismatches 0; Indels
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|||||||
1 FYDPMFK 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TREET: ]
TERR READABLE FORM:
THER THOMPY disk
PUTER: INDAPPY disk
PUTER: IEM 330 466 D.N2
PUTER: IEM 330 466 D.N2
PUTER: PATENTIN: PC-DOS/MS-DOS
PUTER: PATENTIN: PC-DOS/MS-DOS
PUTER: PATENTIN: PC-DOS/MS-DOS
PUTER: PATENTIN: PC-DOS/MS-DOS
PUTER: PC-D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                United States of America
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ON: PROSTATE-SPECIFIC MEMBRANE ANTIGEN AND DR: USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Warren D.W. William R.
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RESULT 29
US-08-946-3129A-82
US-08-946-3129A-82
Sequence 82, Application US/08946329A
PACENT No. 6037991
APPLICANT: Beachy, Philip A.
APPLICANT: Beachy, Philip A.
APPLICANT: Beachy, Philip A.

CANT: Beachy, Philip A.

CANT: POTTER, JEFFREY A.

OF INVENTION: NOVEL HEDGEHOG-DERIVED POLYPEPTIDES

OF INVENTION: NOVEL HEDGEHOG-DERIVED POLYPEPTIDES

REET: TA CA

PONDENCE ADDRESS:
ESSEE: Fish & Richardson P.C.
ET: 4225 Executive Square, Suite 1400

Mon

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RECISPANTON YUDEEN 38.347
REFERENCE/DOCERN WHOREN 07265/140001
TELEPONE 19.5/878.500
INFORMATION FOR SEQ IN NO. 82.
SEQUENCE CHARACTERISPICS
SEQUENCE CHARACTERISPICS
LENGTH: 106 amino acids
TELEPAX amino acids
TELEPAX PER LANGUAGE
LENGTH: 106 amino acids
TELEPAX PER LANGUAGE
NO. 2012 TYPE LORGING PER LANGUAGE
NO. 2014 C. 2014 TYPE LORGING PER LANGUAGE
US-08-946-129A-82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PILING DATE: 07-CCT-1996
APPLICATION UNMBER: 08/729,743
FILING DATE: 10-UUL-1996
APPLICATION WUMBER: 08/567,357
FILING DATE: 04-DCC-1995
APPLICATION WUMBER: 08/39,498
FILING DATE: 07-DFC-1994
APPLICATION WUMBER: 08/39,498
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Query Match 1.0%; Score 7; DB 3; Length 106;
Best Local Similarity 100.0%; Pred. No. 22;
Matches 7; Conservative 0; Mismatches 0; Indels
22 KAENIKK 28
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44 KAENIKK 50
                                                                 0; Indels
                                                                      0,
                                                                      Gaps
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0

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3ULT 30
3-08-970-133-3
"ence 3, Appl
                                        UP INVENTION: SH3-CONTAINING PROTEINS WOMER OF SEQUENCES: 5

OMRESSONDENCE ADDRESS:

ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Dr.
TITT: Palo Alto
TMTE: CA

DUNTRY.
ZIP: 94304
MPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                    Application US/08970133
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Mon Apr 9 11:14:12 2001
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Page 56

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ALL:
AND CONTRY: U...

APPLY 92037

MEMORIAN PRE: Diskette

MEDIUM TYPE: Diskette

MEDIUM STERME TO CONTROLOME

APPLICATION NOMER: 05/061/323
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UPOLOGY Incer
INMEDIATE SOURCE:
LIBBARY: PROSNOTZO
CLONE: 1816529
US-08-970-133-3
                                                                                                                                                                                                                                                                                                                              US-09-031-07A-2
US-09-031-07A-2
Sequence 2, Application US/09053197A
Sequence 10 Application US/09053197A
Sequence 10 Application US/09053197A
APPLICANT: Weiner, Joel H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 1.0%; So
Best Local Similarity 100.0%; p
Matches 7; Conservative 0;
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ANDALA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WITER: IBM Compatible
WAIRING SYSTEM: DOS
WARE: FastSSO for Windows Version 2.0
T APPLICATION DATA:
CATION NUMBER: U/08/977 ...
C. DATE: Filed U/08/977 ...
NSEE: Medien & Carroll, LLP
T: 220 Montgomery Street, Suite 2200
San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOS VERL

LOS Piled Herewith

POLOATION DATA:

TION NUMBER:

ATE:
                                                                                                                                                                           Heiner, Joel H.

WITHER, RAYMOND J.

WITHON: COMPOSITIONS AND METHODS FOR PROTEIN
WITHON: SECRETION
UENCES: 72
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UENCE CHARACTERISTICS
ENGTH: 7 amino acids

PE: amino acid RANDEDNESS: double POLOGY: linear

peptide NO

ORIGINAL SOURCE:
ORANGE MORE ASPLEN
TISSUE MY Home Asplen
TISSUE TYPE: Carcinoma
IMMEDIATE SOURCE:
CLONE: Prostate Specific Membrane Antigen
US-08-394-152A-5

Query Match Similarity 100.0%; Score 7; DB 2; Length 7; Best Local Similarity 100.0%; Pred. No. 1.3+05; Matches 7; Conservative 0; Mismatches 0; Indele

Indels

0;

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521 FYDPMFK 527 |||||||| 1 FYDPMFK 7

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Mon Apr 9 11:14:12 2001
California
Y: United States of America
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                                                                             Page 57
                                                                               Mon Apr 9 11:14:12 2001
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; MOLECULE TYPE: protein US-09-053-197A-2
                                                                                                                                                                                                               Query Match 1.0%; Score 7; DB 3
Best Local Similarity 100.0%; Pred. No. 57;
Matches 7; Conservative 0; Mismatches
698 AAETLSE 704
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231 AAETLSE 237
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AFORMATION:
AFORMA
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RESULT 32
US-088-847B-13
Sequence 13, Application US/OB4ev...
Patent No. 5780263
Patent No. 5780263
CERENT, HERNATIONS, Gregg A. and Adams, Mark D. APPLICANY: Hastings, Gregg A. and Adams, Mark D. 1711E OF INVENTORS; Human CCH-Like Growth Factor UNINESSON BEOGRAPHS.
ONNESSE: OWNER, AND AND GIVETLIAN, GIVENLAN, GIVETLIAN, GIVETLI
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TITILE OF INVENTION: DNA Molecules Encoding Plat
TITILE OF INVENTION: Protoporphyrinogen Oxidase
TITILE OF INVENTION: Thereof
UNDER OF SOURCES: 15
DOBESSORDENCE ADDRESS:
ADDRESSEZ: ADDRESSEZ: DOWNLES PLATE OF SOURCESSEZ: DOWNLESSEZ: 520 White Plains Road, P.O. Box 2005
                                                                        DNA Molecules Encoding Plant
Protoporphyrinogen Oxidase and Inhibitor-Resistant Mutants
Thereof
                                                                                                                                                                                                                                us-08-403-803-2_copy_45_750.olig.rai
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Mob

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TYPE: Ploppy disk
ER: IBM PC compatible
ING SYSTEM: PC-DOS/MS-DOS
ING SYSTEM: PC-DOS/MS-DOS
APPLICATION DATA
APTION NUMBER: US/08/808,931
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ATCAN:
ATION NUMBER.
ATION NUMBER.
ATION NUMBER:
ATION NUMBER:
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ATION THORNATION:
ATRIANTON NUMBER: 30 241
ATTON NUMBER: 30 241
ATTON NUMBER: 30 241
ATTON NUMBER: 30 341
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PPLICATION DATA: US 60/012,705
ANTION NUMBER: US 60/012,705
DATE: 28-FEB-196
PLICATION DATA:
NATION NUMBER: US 60/013,612
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                UMBER: US 60/013,612
28-PEB-1996
CON DATA:
UMBER: US 60/020,003
21-JUN-1996
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Query Match 1.0%; Score 7; DB 2; Length 539; Best Local Similarity 100.0%; Pred. No. 1.1e-0; Matches 7; Conservative 0; Mismatches 0; Indels
102 PPPPGYE 108
                                         0
                                  Gaps
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0;

MOLECULE TYPE:

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Hon
Apr 9 11:14:12 2001
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187 PPPPGYE 193

Page 59

us-08-403-803-2\_copy\_45\_750.olig.rai

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APPLICANT: Johnson, Marie
APPLICANT: Wollath, sandra
APPLICANT: Marie Eric
APPLICANT: Marie Eric
APPLICANT: Marie Eric
APPLICANT: Marie Eric
APPLICANT: Marie
A
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UNTRY: USA

IF: 10541-9005

IF: 10521-9005

MEDIUM TYPE: Floppy disk
COMPUTER: ENDER PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPTRATE: PR-STEM: PC-DOS/MS-DOS
OPTRATE: NEGETIN BLAGE #1.0, Version #1.30
OPTRATE: NEGETIN BATA:
US/08/808,323
NAME: Neigs, J. Timothy
REGISTRATION NUMBER: 38 241
REGISTRATION NUMBER: 36 261
RETERONATION NUMBER: GCC 1846
RELECONDUNICATION LIPORNATION:
TELEPHONE: (919) S41-8689
DRANTION FOR SEG 1D No. 16:
9018NCS CHARACTERISTICS:
LENGTH: 539 anino acids
TYPE: maino acids
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ORAPPICATION 800
OPACADAM NUBERIA US 60/012,705
OPACADAM NUBERIA US 60/013,705
OPACADAM NUBERIA US 60/013,612
PELCATION NUBERIA US 60/013,612
PELCATION NUBERIA US 60/013,612
ORAPPICATION NUBERIA US 60/020,003
ILING DATE: 13-UN-1996
ORAPPICATION NUBERIA US 60/020,003
ILING DATE: 13-UN-1996
ORAPPICATION NUBERIA US 60/020,003
ILING DATE: 13-UN-1996
ORAPPICATION TUBERIA US 60/020,003
ILING DATE: 13-UN-1996
ORREY/ORBET IMPORMATION: US 60/020,003
ORREY/ORBET IMPORMATION: US 60/020,003
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6018105
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Page 58

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ADMPUTER: 138 195/3
COMPUTER: 138 195/3
PILLING DATE: 43.
                                                                                                                                                                                                               STATE: NEW JERSEY
COUNTRY: USA
                                                                                                                                                                                     TER READABLE FORM
                                                                                    GENT INFORMATION:
GULLINS, J.G.
TION NUMBER: 33,0
                                                                            325800-442
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ouery HAtch 10.0%; Score 7; DB 1; Length 375; Best Local Similarity 100.0%; Pred. No. 75; Best Local Similarity 00.0%; Simmatches 0; Indels Matches 7; Conservative 0; Mismatches 0; Indels
                                                                            MOLECULE TYPE: PROTEIN -08-468-847B-13
         0
      Gaps
```

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ESUIL 3
15-08-808-931-16
Sequence 16, Application US/08808931
Patent No. 593960
GENERAL INFORMATION:
APPLICANT: Volrath, Sandra
                                                                                                                                                                           454 TKKSPSP 460
|||||||
284 TKKSPSP 290
Volrath, Sandra
Johnson, Marie
Potter, Sharon
Ward, Eric
Heifetz, Peter
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DB 3; Leng No. 1.1e+02; No. 1.ndels

Gaps

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US-09-050-603A-16
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TOPOLOGY: not relevant
: MOLECULE TYPE: protein
US-08-808-323-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ouery Match 10.0%; Score 7; DB 3; Length 539; Best Local Similarity 100.0%; Pred No. 1.1e+02; Hatches 7; Conservative 0; Hismatches 0; Indels
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|||||||
187 PPPPGYE 193
                                    AGENT INFORMATION MARGEST AND TAMOSTY AND TAMOSTY AND TAMOSTY AND TAMOSTY NOMER: 35,241
MACINORMATION WHERE: 36,241
MACINORMATION THOREM: 360,741
MACE CHARACTERISTICS:
ACE CHARACTERISTICS:
GOTH: 539 amino acids
GOTH: 539 amino acids
GOTH: 539 amino acids
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25SEE: No. 602301artis Corporation
77: 3054 Cornwallis Road
Research Triangle Park
1. NC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ( REMANDLE PORH:

( TTPE: Ploppy disk

(TER PC oppy disk

(TER ISH PC ompatible

ING STSTEM: PC-DSS/MS-DGS

ARE: PAtentin Release #1.0, Version #1.30

ARE: NEW TOWN NOTA:

TE /NO /NGO. 603A
                                                                                                                                                                                                                                                                                                                                                                                          DATE: 06-JUN-1995
AGENT INFORMATION
TELSS, J. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                          LLANIU US 60/020,003
LLANIU US 60/020,003
LLANIUN DATA:
"ION DATA: US "IS"
"ION BURBER: US "IN"
PAIR OF "
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DATE: 30-9AR-1299/050,603A
DATE: 30-9AR-1299
TICATION DATE: US-08/008,931
DATE: 28-FEB-1997
ULCATION DATE: US-60/012,705
DATE: 28-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Volrath, Sandra
Johnson, Marie
Potter, Sharon
Mard, Eric
Heifett, Peter
WestTION: DNA tolecules Encoding Plant
WestTION: Protoporphyrinogen Oxidase
ESQUEREES: J7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28: US 60/013,612
28:FEB-1996
VON DATA:
UMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TI-MAR 1998
N DATA: UBER:
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M PC compatible

NIEM: PC-DS/MS-DOS

TIEM: PC-DS/MS-DOS

(TION DATA: 05.09/102,4208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ER: US 60/012,705
20-FEB-1996
N DATA:
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MABER: US 09/059,164

13-APR-1998

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M DATA:

MBER: US 09/050,603

30-MAR-1998

30-MAR-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mon Apr 9 11:14:12 2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches 7; Conservative
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Best Local Similarity 100.0%; Pred. No. 1.1
Matches 7; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTOR: A METHOD FOR DETECTING A MANMAL'S PRIOR TITLE OF INVENTOR: A METHOD FOR DETECTING A MANMAL'S PRIOR TITLE OF INVENTOR: EXPOSURE TO RADIATION OR RADIOMIMETIC AGENTS UNDERS OF SEQUENCES: 2

CORRESPONDENCE ADDRESS;
ADDRESSEE: HANOR, HATSTRAMP, Devans 6 Doyle LLP
STREET: CLINION SQUARE, P.O. BOX 1051

STREET: CLINION SQUARE, P.O. BOX 1051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         102 PPPPGYE 108
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187 PPPPGYE 193
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187 PPPPGYE 193
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ER READABLE PORM:
UM TIPE: Floppy disk
UTER: IBM PC compatible
OUTER: IBM PC compatible
ANTING STSTEM: PC-DOS/MS-DOS
HARE: PAtentin Release #1.0, Version #1.30
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MMUNICATION INFORMATION:
PHONE: (919) 541-8587
PHONE: (919) 541-8689
INFORMATION: 1619
INFORMATION: 1619
INFORMATION: 1619
INFORMATION ACIDS
INFORMATION
INFORM
                               ... AUMBER: __3,601
MCATOM INFORMATION:
NE. (716) 263-1634
(716) 263-1630
FOR SED ID NO. 1:
**ARACTERISTICS:
622 anino acids
ino acid
**Course | Course | Co
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1054 Cornwallis Road
Search Triangle Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FION DATA:
JMBER: US/08/547,197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                J. Timothy
NUMBER: 38,241
CGC 1847
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28-FEB-1996
ON DATA:
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: OXIDASE ("PROTOX")
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0; Indels 0; Gaps

Page 64

Query Match 1.0%: Score 7; DB 3; Length 539; Best Local Similarity 100.0%; Pred. No. 1.1e+02;

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AER OF SEQUENCES:

JRRESPONDENCE ADDRESS:

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US-08-959-40-1
Sequence 1, Application US/08957940
Patent No. 6132981
Patent No. 6132981
GENERAL INFORMATION:
GENERAL INFORMATION:
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|||||||
190 GLPSIPV 196
                                                                       TORNET/AGENT INPORMATION:

MAE: Kognlakyj, Peter

REGISTRATION NUMBER: 39,601

REGISTRATION NUMBER: 19226/580 (R-5228)

RECHENNEZ/DOKERT NUMBER: 19226/580 (R-5228)

RECHENNEZ/DOKERT NUMBER: 19226/580

RELEPAX: (715) 261-1600

AATION FOR ESD ID NO: 1:

RECHE LURACTERISTICS:

NOTH: 672 Amino acida

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: A METHOD FOR DETECTING A MANMAL'S PRIOR
:: EXPOSURE TO RADIATION OR RADIOMINETIC AGENTS
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US-08-325-553-35
US-08-325-553-35
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Best Local Similarity 100.0%;
Matches 7; Conservative
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190 GLPSIPV 196
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ASSIFICATION 435
NA ADRICATION MINISER: US/07/973,337A
PLIANTON MINISER: US/07/973,337A
PLIANTON MINISER: US/07/973,337A
PLIANTON MINISER: 14/4/4426
PERSONS/PLATION MINISER: 14/4/4426
PERSONS/PLATION MINISER: 14/4/4426
PERSONS/PLATION MINISER: 12/4/44426
PERSONS/PLATION MINISTER STORE
PERSONS (22) 16/4-0325
PERSON
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PMOVIES: IBM PC compatible
BRATING SYSTEE: PC-DOS/MS:DOS
FTMARE: Patentin Release #1:0, Version #1:25
ENT APPLICATION NUMBER: US/08/325,553
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Heston, Marren D.M.
Reit, Hilliam R.
Reminon: THE PROSTATE SPECIFIC MEMBRANE ANTIGEN
GUENCES: 38
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; Pred. No. 1.2e+02;
0; Mismatches 0; Indels
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ORGANISM, Homo Sapian
CELL TYPE, Carcinoma
CILLE, POBETIATE SOURCE:
CLORE: POBETIATE SPECIFIC Membrane Antigen
US-08-325-553-36
                                                        Apr 9 11:14:12 2001
                                                                                                                                                                                                                                                                 Query Match

Best Local Similarity 100.0%; Score 6; DB 1; Length 6;
Best Local 6; Conservative 0; Mismatches 0; Indels
Matches 6; Conservative 0; Mismatches 0; Indels
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||||||
1 NEDGNE 6
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CULE TYPE: peptid-
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SERENCE/DOCET NUMBER: 1747/41426
SHENCE/DOCET NUMBER: 1210/2550
SHOWNICATION INFORMATION I
SPAN: (212) 564-5555
SPAN: (212) 564-555
SPAN: (212) 565-555
TOM TORS SEGL DU NO. 36:
ESSEE: Cooper & Dunham
ET: 30 Rockefeller Plaza
: New York
: New York
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CTERISTICS
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RESULT 40 US-08-225-533-36 IS-equence 36, Application US/0832553 Retent No. 553886 PATCH INFORMATION:

PLICANT: Heaten Marren D. M.
PLICANT: Heaten Marren D. M.
PLICANT: PAIT, WILLIAM R.
PLE OF INVENTOR: THE PROSTATE-SPECIFIC MEMBRANE ANTIGEN
DES OF SEQUENCES: 39
RESPONDENCE ADDRESS:
POPRESSEE: Cooper & Dunham
REET: 30 ROCKET-LIET Plaza
TT: New York
ATE: New York
ATE: New York

CONTROL OF THE PROPERTY OF THE

New York E: New York TRY: United States of America

Mon

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9 11:14:12 2001

us-08-403-803-2\_copy\_45\_750.olig.rai

Page 68

STRANDEDNESS: single TOPOLOGY: linear LECULE TYPE: protein

double linear linear TYPE: peptide TTICAL: NO SE: NO SOUTH

TH: 6 amino acids : amino acid NDEDNESS: double

Query Match 0.8%; Score 6; DB 1; Length 6; Beat Local Similarity 100.0%; Pred, Mo. 1.3++05; Matches 6; Conservative 0; Mismatches 0; Indels

0 Gaps

19 DELKAE 24 |||||| 1 DELKAE 6

United States of America

eli, Ron S. on, Warren D.W. William R.

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JOBER: 19,

JOHN 1992

JAT INFORMATION

JATON NOMBER: 13,47/41426

MUCLOCACET NUMBER: 1747/41426

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MUCLOCACET NUMBER: 1747/41426

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MUCLOCACET NUMBER: 17,17
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ORAMISH: Bomo Sapien
CELL TYPE: Carcinoma
INEDIATE SOURCE:
CLONE: POOSTATE Specific Membrane Antigen
US-08-325-553-37
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US-08-394-152A-35
; Sequence 35, Application US/08394152A
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                                                                                                                                                                                                                                                                 RECISTRATION NUMBER: 28.678
REFERENCE/DOCKET NUMBER: 4126-B
TELEPONERICATION IMPORMATION
TELEPHONE: (212) 278-0400
TELEPHONE: (212) 278-0400
TELEPHONE: (212) 291-0525
TELEPHONE: (316) 391-0525
TELEPHONE: (316) 391-0525
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-394-152A-36
uence 36, Application US/08394152A
ent No. 5935818
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1 DELKAE 6
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PILCANT: PAIT, WIlliam R.
THE OF INVENTION: PROSTATE-SPECIFIC MEMBRANE ANTIGEN AND
THE OF INVENTION: USES THEREOF
WEER OF SEQUENCES: 48
RESPONDENCE ANTERIOR
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DDNESSEE: Cooper & Dunham LLP
DDNESSEE: 1185 Avenue of the Americas
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ving System: PC-pos/w*

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R: IBH 304 646 DX2
R: IBH 304 646 DX2
R: SYSTAN OC-DOS/MS-DOS
BE Detenth Release $1.0, Version $1.25
BE DELCATION DATA:
FINAN UNITED NATA:
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ICATION DATA:
ION NUMBER: US/07/973,337A
ATE: 05 NOV 1992
JENT INFORMATION:
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ICATION DATA:
N NUMBER: US
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Release #1.0, Version #1.25
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Pred. No. 1.3e+O5;
0; Mismatches 0; Indels
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: IMMEDIATE SOURCE:

: CLONE: Prostate Specific Membrane Antigen
US-08-394-152A-36
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CLONE: Prostate Specific Membrane Antigen
US-08-394-152A-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.8%; So
Best Local Similarity 100.0%; I
Matches 6; Conservative 0;
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08-394-152A-37
08-394-152A-37
Acquence 37, Application US/08394152A
Actent No. 593588
Actent No. 5935818
Actent No. 5935888
Actent No. 5935888
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1 NEDGNE 6
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YTONBET/ACERT INFORMATION:
NAME: WHICE, John P.
REGISTRATION: NUMBER: 28.678
REFERENCE/COCKET NUMBER: 41426-B
RELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 278-0400
TELEPHONE: (212) 29-0525
ORDANION FOR SEQ ID NO: 37:
SEQUENCE CHARTON FOR SEQ ID NO: 37:
                                                                                                                                                                                                                                                                                          NEUTIER READABLE FORM:
MEDIUM TYPE, PLOPPY disk
MEDIUM TYPE, PLOPPY disk
ORMUTER: IIBM 330 466 DN2
OPERATIR STETEM: PC-DOS/MS-DOS
OPERATIR STETEM: PC-DOS/MS-DOS
OPERATIR DATE:
MARBUT APPLICATION DATM:
MAPPLICATION MORRE:
05/08/394,152A
PPLICATION MORRE:
03/08/394,152A
PLING DATE: 24-PEB-95
CLASSIPICATION: 435
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ITEM TYPE: Floppy disk
POTER: 18 330 466 DX2
RANING SYSTEM: DC DOS/MS-DOS
THARE: Patentin Release #10, Version #1.25
NT APPLICATION DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: Carcinoma
ATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ONDENCE ADDRESS:
SSEE: Cooper & Dunham LLP
ST: 1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            )GY: linear
! TYPE: pep
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VENTION: PROSTATE-SPECIFIC MEMBRANE ANTIGEN AND VENTION: USES THEREOF
EQUENCES: 48
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United States of America
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Cooper & Dunham LLP

Don't he Americas
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United States of America
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24-FEB-95
: 435
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USES THEREOF
5: 48
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b. 1.3e+05;
ches 0; Indels
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Mon Apr 9 11:14:12 2001
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ORGANISH: Homo Sapten
CELL Type: Carcinoma
IMBELIATE SOURCE: CONTROL CLONE: Protekte Specific Membrane Antigen
US-08-194-152A-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query March 0.8%; Score 6; DB 2; Length 6;
Beat Local Similarity 100.0%; Pred, No. 1.3+05;
Matches 6; Conservative 0; Mismatches 0; Indel:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       438 KSPDEG 443
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1 KSPDEG 6
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mence 12, Application US/07968781A
mt No. 5430137
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R READABLE FORM:
A TYPE: Floppy disk
TETK_IBH PC compatible
TRING SYSTEM: PC-DoS/MS-DOS
ARE: PAtenting Release #1.0, Version #1.25
APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          H: 6 amino acids
amino acid
DEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     David R. Saliwanchik
1421 N.W. 41st Street, Suite A-1
nesville
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b. George E.
, Kanneth E.
W: Probes for the Identification of Bacillus
W: thuringieneis Endotoxin Genes
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PCT-0821-10.68 4
RCT-0822-10.68 4
RCT-0822-10.68 4
RESULTANT RESULTANT DATE OF APPLICANT Edgington, Thomas S
APPLICANT Edgington, Thomas S
RESULTANT RESULTANT PROFILE OF THE RESULT NO. 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-68 10-6
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Best Local Similarity 100.0%;
Matches 6; Conservative 0;
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1 GPGFTG 6
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PORMATION FOR SEQ ID NO:
PORMATION FOR SEQUENCY IN SEQ ID NO:
PORMATION FOR SEQ ID NO:
PO
       MPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PPLICATION NUMBER: US/07/968,781A
ILING DATE: 19921030
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OLE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                      Office of Patent Counsel, The Scripps: Research Institute
10666 North Torrey Pines Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ENTION: Factor X-Derived Polypeptides and Therapeutic Methods ENTION: Anti-Peptide Antibodies, Systems and Therapeutic Methods ENTION: for Inhibiting Inflammation QUENCES: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12:
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OPERATING SYSTEM: PC-DOS/MS-DOS
CURRENT PARTIC TION PARTIC STORY
CURRENT PARTIC TION PARTIC STORY
CLASSIFICATION NUMBER: PCT/USS2/10068
FILING DATE: NUMBER: PCT/USS2/10068
FILING DATE: 22-MSV-191
FILING DATE: 22-MSV-191
FILING DATE: 22-MSV-191
FILING DATE: 1519-554-291
FILING DATE: FIRETING HURBER: SCR1221P
REFERENCE/DOCKET NUMBER: 34,63
FELECOMMUNICATION HURBER: 34,63
FELECOMMUN
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Mon Apr

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Page 75

Mon

Apr 9 11:14:12 2001

us-08-403-803-2\_copy\_45\_750.olig.rai

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APPLICANT, Gaeta, Redariko
APPLICANTO, Latander, Jesthalin Peptides
TITLE OF INVENTION: Destander Peptides
NUMBER OF SEQUENCES: 39
CORRESTE STEED APPLICATION APPLICATION APPLICATION APPLICATION APPLICATION MATA:
APPLICATION MUMBER: US 08/121/101
TELEPHONE: (145) 576-0300
TELEPHONE: (145) 576-0300
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TYPE: ABIND CAIGS: 11 ABIND CAIGS
TYPE: ABIND CAIGS
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Page 77

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Page 78

; MOLECULE TYPE: peptide US-08-164-618-1

Ouery Match 0.0%; Score 6; DB 1; Length 15; Best Local Similarity 100.0%; Pred, No. 36; Matches 6; Conservative 0; Mismatches 0; Indels

0; Indels

0

Gaps 0

258 AQKLLE 263 ||||||| 3 AQKLLE 8

RESULT 49
US-08-156-552A-6
Sequence 6, Application US/08156552A
Patent No. 5726155
Patent No. 5726155
GENERAL INFORMATION:

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Best Local Similarity 100.0%; Pred. No. 32; Matches 6; Conservative 0; Mismatches
697 AAAETL 702
|||||||
4 AAAETL 9
                                                 0;
                                                 Indels
                                                 Gaps
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RESULT 48
US-08-164-618-1
US-08-164-618-1
JSequence 1, Application US/08164618
JSEQUENCE 1, Application US/08164618
JEREAL INFORMATION:
GRANITON:
APLICATION WALL DEPTH 1988
FILIKATIES 1990
APTORNEY JGENT INFORMATION:
NAME: GAMENO EMPACE
REGISTRATION NUMBER: 29.381
REFERENCE/DOCKET WUMBER: SCRF 231.0
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATICS:
LEWRIE: 13 amino acida
TOPOLOCE: LIABRIE AMINO acida
TOPOLOCE: LIABRIE AMINO acida
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOVIEW POODS.

MOVIEW THE MAD POORM

MEDIUM THE MAD POORM

COMPUTER: INW POOPWHARTING

OPERATING SYSTEM: PO-DOS-MOS-MOS-

OPERATING SYSTEM: PO-DOS-MOS-MOS-

OPERATING SYSTEM: PO-DOS-MOS-MOS-

NOTWHARE APPLICATION DATA:

NET APPLICATION NUMBER: US/08/164,618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DDDBESSEE: Treeslar Goldsmith, Shore, Sutker & ADDBESSEE: Willman, Ltd. Stetaon, Suite 4700 STREET: 180 No. 3408036th Stetaon, Suite 4700 STREE: Chicago.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LICANT: Chadiri, M. Reza
LIZ OF INVENTION: Isolated Metallopolypeptides:
LE OF INVENTION: Compositions and Synthetic Methods
BER OF SEQUENCES: 20
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LICATION DATA:
TION NUMBER: US/07/769,621
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RESULT 50
US-08-477-509B-112
Sequence 112, Application US/08477509B
Patent No. 5770697
GENERAL IMPORMATION:
APPLICANT: Ferrari, Franco A
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PRICANT: Capello, Joseph

PRICANT: Capello, Mary

PRICANT: Capello, Joseph

PRICANT: Capello, Joseph

PRICANT: Capello, Joseph

ADDRESSEE: Paler Description

PRICANT: San Francisco

ETMTE: California
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Indels
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Best Local Similarity 100.0%; Pred: No. 39;
Matches 6; Conservative 0; Mismatches 0; Indels
                                                                                          NOLECULE TYPE: protein :-08-156-552A-6
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258 AQKLLE 263
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3 AQKLLE 8
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SEQUENCE CHARACTERISTICS: LENGTH: 16 amino acids TYPE: amino acid TOPOLOGY: linear

NEDUTER REALMANCE PORM.

MEDIUM TIPE: PLOPPY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMANE: Patentin Re-Lease #1.0, Version #1.30
MPDICATION NUMBER: US/08/477,509B
FILING DATE: 07-UN-1995
CLASSIFICATION: 455

Mon Apr 9 11:14:12 2001

APPLICATION DATA:
URATION NUMBER: US 08/175,155
NG DATE: 29-DEC-1993
NFPLICATION DATA:
UCATION NUMBER: US 08/053,049
NG DATE: 22-AFR-1993

Page 79

Mon

Apr 9 11:14:12 2001

us-08-403-803-2\_copy\_45\_750.olig.rai

TELEFAX: 619-554-6312 INFORMATION FOR SEQ ID NO:

APPLICATION NUMBER: "US/08/156,552A FILING DATE: 15-MOV-1993 PRICESSIFICATION: 5.14 PRICESSIFICATION: 5.14 PRICESSIFICATION: 5.14 PRICESSIFICATION: MATRIX US 08/102,944 PTILING DATE: 02-ANG-1993 VITONBENE/AGENT INFORMATION: MATRIX LOSAN, APTIL C. REGISTRATION NUMBER: 30.950 REFERENCE/DOCKET NUMBER: SCRF 281.1 PRICESSAL 2837 PRICESSAL

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TIZE OF INVENTION: REGULATION OF OXIDATIVE BURST USING TIZE OF INVENTION: LAWG-DERIVED PETTIES AND ANLOGS MEER OF SEQUENCES: 31
RESEPONDENCE ADDRESS:
RESEPONDENCE ADDRESS:
Factor Counsel Institute, Office of ADDRESSEE: Acta Counsel Tourier Pines Road, TPC 8
STREET: 10666 No. 5726155th Torrey Pines Road, TPC 8
STATE: La Jolla
STATE: USA

BOKOCH, GARY M
CUTNUTTE, JOHN
FENTION : GROULATION OF OXIDATIVE BURST USING
FENTION: LANG-DERIVED PEPTIDES AND ANALOGS
QUENCES: 31
FE ADDRESS: 31

APPLICATION NUMBER: 29-OCT-1987 N DATA:

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RELECOMMUNICATION THE CAMATION:
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STRANDEDURES: single
VOLECULE TYPE: Innat
VOLECULE TYPE: BER: US 06/927,258

STRANDEDNESS: single;
TOPOLOGY: linear;
MOLECULE TYPE: peptide
US-08-477-509B-112

Query Match
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Best Local Similarity 100.0%; Score 6; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 50;
Matches 6; Conservative 0; Mismatches 0; Indels Indels o, Gaps

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Search completed: April 6, 2001, 16:56:59 Job time: 133 sec

s the number of results predicted by chance to have a printed, than or equal to the score of the result being printed, and by analysis of the total score distribution.

SUMMARIES

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on:

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- protein search, using sw model

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MESSON; the aproactate specific membrane antigen (PBM, encoded by 065250). The inventors have identified a peptides of high hydrophilicity to which antibodies may be directed against. In the specific membrane spanning domain as stated to be from anion cold residue 19 to residue 44. The PBM coding sequence is useful for acid residue 19 to residue 44. The PBM coding sequence is useful for appreciation or modulating the metastic billity of prostate tumour cells to grow, or for eliminating them. The protein is useful to identify or purify ligands of the Ag It is also an attractive target for Ab-directed imaging and targeting of prostatio unour deposits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 24; Page 103-106; 196pp; English.
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423 dctplmyelvhnitkelkepdegfegkelyeevtkkepepefegmprisklydgndfevf 482
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225 vilysdradfrafydrafyddymilpsgyraininingagdpilpsgypancysyrigia 204
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165 spogmpsgolvyvnyaftenpfilendmingskivtarygkvfignkvhnaqlagakg 224
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63 ldavelahydvilayphkyhphyisiinedgnoiintaifeppppyyenvadivppfsaf 122
                               361 LKEGHRPRETIFASHVAREPGIJGSTEWARENSHLIGERGVATHANDSIEGREVTILRV 420
405 LKRGGVEPFITLITASHVAREPGIJGSTEWARENSHLIGERGVATHANDSIEGREVTILRV 464
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105 ldavelahydvllaypnkthepnylsilnedgneifntslfeppppgyenvadivppfsaf 164
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Y41074;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Y41074 standard; protein; 750
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N-PSDB; Q65520.
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                                                                                                                      Murphy GP, Boynton AL, Holmes EH, Tino WT
                                                                                                                                                                                                                                                                                                   18-MAR-1999;
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PSMA; prostate specific membrane antigen; PSM' protein; prostate cancer.
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                                                           WPI: 1999-580294/49.
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481 PORIGIASGARIYTKANETMKFSGYPLYBSYETYELVEKETVDPKEKHLTVAVGGGW
525 fqr1giasgrarytknwetnkfsgyplyhavyetyelvekfydpmfkyhltvaqvrggmv
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185 apgunpegdivyvoyartedtrierdmkincagkiviarygvutgykkanolistasasg 224
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    /label= insertion_region
/note= "suitable for foreign T helper epitope insertion"
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/note= "computer-predicted antigenic determinant"
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                                           /label= antibody_binding_region
/note= "computer-predicted antigenic determinant=
442..465
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/note= "suitable for foreign T helper epitope insertion"
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/note= "experimentally found B cell epitope"
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= "subdominant CTL epitope"
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        Claim 56; Page 184-187; 220pp; English
                                Inducing immune responses to weakly immunogenic, tumor associated peptide antigens for the treatment of breast and prostate cancer
                                                                         WPI; 2000-349917/30
N-PSDB; A09454.
                                                                                                              Steinaa L. Mouritsen S. Nielsen KG. Haaning J. Leach D. Dalum Gautam A. Birk P. Karlsson G;
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20-0CT-1998;
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98US-0105011
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/label= insertion_region
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58..750
/label= PSM'
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/note= "subdominant CTL epitope"
/16..723
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note= "experimentally found B cell epitope and
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465 dctplmyslvhnltkelkspdcsfcgkslyesvtkkspspefsgmprisklgsgndfevf
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                                                    05-OCT-1999;
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17.31
/label= P2
/note= "foreign epitope"
32.52
/label= P30
/note= "foreign epitope"
                       98DK-0001261
                                                    99WO-DK00525
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24..38
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Pred. No. 0;
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0; Indels Length 750;

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XX XX XX Y2627-49 are mutant immunogenized human prostate specific membrane CC y2627-49 are mutant immunogenized human forzign epitopes (P2 and/or P30). CC antigen (PBM) constructs, which contrain forzign epitopes (P2 and/or P30). CC The immunogenic analogues of PBM can be used in the claimed method as an attractive to the process of the contraint of the contraint of the process of the contraint of the contraint of the process of the contraint of the contraint of the process of the contraint of the contraint of the process, e.g. thuman prostate specific membrane antigen (PBM), hereguin 2 (Ber2) and/or CC immune system of: (1) at least 1 CT. (cytocota: Tlumphocyte) contraints at least 1 (TT. (cytocota: Tlumphocyte) group derived from the PA and/or at least 1 (TT. (cytocota: Tlumphocyte) group derived from the PA and/or at least 1 (TT. (cytocota: Tlumphocyte) group derived from the PA and/or at least 1 (TT. (cytocota: Tlumphocyte) group derived from the PA and/or at least 1 (TT. (cytocota: Tlumphocyte) group derived from the PA and/or at least 1 (TT. (cytocota: Tlumphocyte) group derived from the PA and/or at least 1 (TT. (cytocota: Tlumphocyte) group derived from the PA and/or at least 1 (TT. (cytocota: Tlumphocyte) group derived from the PA and/or at least 1 (TT. (cytocota: Tlumphocyte) group derived from the PA (TT. cytocota: Tlumphocyte) group derived from the path of the apecification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Inducing immune responses to weakly immunogenic, tumor associated peptide antigens for the treatment of breast and prostate cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Steinaa L, Mouritsen S, Nielsen RG,
Gautam A, Birk P, Karlsson G;
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Example 1: Page -: 220pp: English.  XX This is a splice variant of human prostate specific membrane antigen CC [784] (see 192619). Imminogenic analogues of PSM can be used in the CC [784] (see 192619). Imminogenic analogues of PSM can be used in the CC claiminogenic call-and yluthing regions and ystitus expectives in the CC disulfide bonds are preserved in the immunogenized forms. Regions continued for the intertion of foreign T helper spicops were identified. CC immunogenic call-anacolated speciale antigens (PA) such as those continuities and the prostate of the method is used for inducing immunogeness egainst weakly considered with owners (i.e. self-prostatina). For example, human regions of the anacolated speciale antigens (PA) such as those continuities of the self-anacolated speciale antigens (PA) such as those continuities of the self-anacolated speciale and producing cells (APAS) of the call-anacolated place of the self-analogues of human producing cells (APAS) of the call-anacolated place of the self-analogues of human producing cells (APAS) of the call-anacolated place of the self-analogues of human producing cells group bricks to creign to the enimal. Analogues of human producing cells group bricks to creign to the enimal. Analogues of human producing cells group which is contained to the prostate of the self-analogues of human producing cells group which is contained to the prostate of the producing cells group which is contained to the prostate of the producing cells group which is contained to the prostate of the producing cells group which is contained to the prostate of the producing cells group which is contained to the prostate of the producing cells group which is contained to the prostate prostate producing cells group which is contained to treat prostate prostate producing cells group which is contained to the prostate prostate producing cells group which is contained to the prostate prostate producing cells group which is contained to the prostate prostate producing cells group which is con	Db 173 dlyynnyartedffilerdmkincsgkivlarygkvfrynkvhanqlagakgvilyadpa 232  97 189 DPPANOTSSPDORHLEGGOVGROHILHJAGAGDTJPSPANETANRGLISE 248  Db 211 dyfagyfwlydedylleggovycynilhlagodplipypanoyaytrylaeawylpai 292  97 249 PWPIGTONGKILEZKGGGAPPDGSRRSLKYPKNGGFTANESGAVOHINSTNEW 308  Db 331 DYDJSYNAGALLEZKGGGAPPDGSRRSLKYPKNGGFTANESGAVOHINSTNEW 309  98 331 DYDJSYNAGALLEZKGGGAPPDGSRRSLLOBGGANANGENSTRATTSPANEW 309  99 315 DYDJSYNAGALLEZKGGGAPPDGSRRSLLOBGGANANGENSTRATTSPANEW 309  90 411 HILLIHLIHLIHLIHLIHLIHLIHLIHLIHLIHLIHLIHL	Mon Apr 9 11:14:10 2001 us-08-403-803-2_copy_45_750.olig.reg
	Page 43	Page 41
Oy 314 VIGTIKGAVEDDRYTILGEHBDSWFGGIDNGGGANVHEIVRBSPGILKEDSWRPRRTIL 373  Oy 174 PASHDAEREGLIGSTENAEDSKAPTGGDDAGGGANVHEIVRBSPGILKEDSWRPRRTIL 373  DD 301 VIGITISH	RW Product cancer: cell-associated peptide entigen: foreign epitope.  XX Romo aspiens.  XX Romo aspien	Mon Apr 9 ll:14:10 2001 us-08-403-803-2_copy_45_750.olig.rag
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Prostate-specific membrane antigen.  Note: Prostate-specific membrane (PNN) antigen (NO2234) is a type II	Mon Apr 9 11:14:10 2001 us-08-403-803-2_copy_45_750.olig.rag	Presion 716723 701	Mon Apr 9 11:14:10 2001 us-08-403-803-2_copy_45_750.olig.rag
	Page 47		Page 45
CC integral membrane pycoprotein that is highly expressed in prostatic tumours and metastadee and is a larget entity for the pyclic and provides an attractive cell surface epicope for antibody effected coll provides an attractive cell aurace epicope for antibody effected coll provides an attractive cell aurace epicope for antibody effected coll and acqueroes as deduced from a colk close (7585) obtd. from coll acqueroe was deduced from a colk close (7585) obtd. from coll acqueroe was deduced from a colk close (7585) obtd. from coll acqueroe was deduced from a colk close (7585) obtd. from coll acqueroe pycloselic. Deen of papers in the first 57 aniso acids of pseudosty cycloselic. Deen of papers in the first 57 aniso acids of pseudosty close for an increase protectic tumourgoression pseudosty in the deminant collection in princip protectic tumourgoression. Pseudosty close for an increase protectic tumourgoression pseudosty in index can be considered in protectic tumourgoression. Pseudosty close for a collection of pseudosty collection of pseudosty collection. Pseudosty close for a collection of pseudosty collection of pseudosty collection. Pseudosty collection of pseudosty collection of pseudosty collection of pseudosty collection. Pseudosty collection of pseudosty collection of pseudosty collection. Pseudosty collec	Mon Apr 9 11:14:10 2001 us-08-403-803-2_copy_45_750.olig.zag	98 NYISIINEDGNEIPPPSLEEPPPPOYENVALADDIVPPRANCESDIJVPWANNEED 141 127 NYISIINEDGNEIPPTSLEEPPPPYNNOALAGAKVILISSPADDFPRACVKSYPGG 201 128 TRIEBDMINSGITVIREGGVERNIVENAQLAGAKVILISSPADDFPRACVKSYPGG 201 129 142 FRIEBDMINSGITVIREGGVERNIVENAQLAGAKVILISSPADDFPRACVKSYPGG 201 120 MILJOGGVAPDGSMCSLAVPIPOYDRIFATROCARAVULESTRUPTSLYNIGITAGA 221 120 247 MILJOGGVAPDGSMCSLAVPIPOYDRIFATROCARAVULESTRUPTSLYNIGITAGA 221 120 247 MILJOGGAPDGSMCSLAVPIPOYDRIFATROCARAVULESTRUPTSLYNIGITAGA 221 121 121 121 121 121 121 121 121 121	Mon Apr 9 11:14:10 2001 us-08-403-803-2_copy_45_750.olig.rag
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21.41
/label p30
/note= foreign epitope*
91.105
/label p2
/note= foreign epitope*
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980S-0105011
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Location/Qualifiers
24..38
/label= P2
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                          В
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Contain products (TL and production) are also claimed. The method is used to treat proporately prostate prostate products of the specific products (PSM, FGF9B and Her2, respectively.
                                                        W 92627-49 are mutant immunogenized human prostate specific membrane contigen (FBM) constructs which contains the propensive sendors 30). The figen (FBM) constructs which contains the propensive sendors and as an examination of the propensive sendors and the sendors are successful to the first sendors are successful to the immunogenized forms. The method is used for inducing community for the proteins to the cancer (FBM), heregulin 2 (Merz) and/or contigent sporth factors b) (FBM), heregulin 2 (Merz) and/or contigent sporth factors b) (FBM), heregulin 2 (Merz) and/or continuals immune system of: (1) at least 1 (TL (systoxist 7-lymphocyte) (2 group derived from the pa and/or at least 1 (TL (systoxist 7-lymphocyte) (2 group derived from the pa and/or at least 1 (TL (systoxist 7-lymphocyte) (2 group derived from the pa and/or at least 1 (TL (systoxist 7-lymphocyte) (2 group derived from the pa and/or at least 1 (TL (systoxist 7-lymphocyte) (2 group derived from the pa and/or at least 1 (TL (systoxist 7-lymphocyte)) (3 group derived from the pa and (2) at least 1 (TL (systoxist 7-lymphocyte)) (3 group derived from the pa and (2) at least 1 (TL (systoxist 7-lymphocyte)) (3 group derived from the pa and part (1) at lation and (2) at least 1 (TL (systoxist 7-lymphocyte)) (3 group derived from the part (1) at lation and (2) at least 1 (TL (systoxist 7-lymphocyte)) (3 group derived from the pa and (2) at least 1 (TL (systoxist 7-lymphocyte)) (3 group derived from the part (1) at lation and (2) at least 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Steinaa L, Mouritsen S, Nielsen KG, Haaning J, Leach D, Dalum I; Gautam A, Birk P, Karlsson G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-OCT-1998;
20-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Inducing immune responses to weakly immunogenic, peptide antigens for the treatment of breast and
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98US-0105011.
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87..107
/label= p30
/note= "foreign epitope"
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cc antigens (PA) such as those associated with cancers (self-process), e.g.
chuman prostate specific membrane antigen (PBM), heregulin ? (Ber2) and/or
cc fibroblast growth factor Bb (FORDb). The method comprises effecting
cc simultaneous presentation by antigen producing cells (ARCs) of the
antimals immune system of: (1) at least 1 CT (cytocoxic Tlymphocyre)
cgroup derived from the PA and/or at least 1 B-cell group derived from the
cc cell-associated PA; and (2) at least 1 first Thelper cell group which is
consign to the animal. Analogues of human bsy, human Her2 and
comman/murler PCPBs comprising a substantial part of all known and
cc human/murler PCPBs comprising a substantial part of all known and
cc human/murler PCPBs comprising a substantial part of all known and
cc human/murler PCPBs comprising a bustantial part of all known and
cc human bsy and the plicopes of the respective PA and including at
company the comprising a bustantial part of all known the company
cc (Y92619), which appears on pages 184-187 of the specification.
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Best Local Similarity 100.0%; Pred. NB. 0;
Matches 629; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; P
Matches 643; Conservative 0;
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1161 SPOGMPBEDLYYVMARTEDFFKLERDMITKSSKIVTARYGKYPRGHKVKHAQJAGAKG
1161 SPOGMPBEDLYVVMARTEDFFKLERDMITKSSKIVTARYGKYPRGHKVKHAQJAGAKG
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105 ldsvelahyovLsyphkthpnylsiinedgneifntsifeppppgyervedivepfsaf 164
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| 164 GHPEGLIYYUKAKTEDFFKLERUMKINGSGKIYLAYGKTVFRGUKVKNAQLAGANGVIL 183
| 168 GMPEGGGIVYUNJAKTGGKKINGSGKIVIATYGKVITGKYKNAGLAGANGVIL 227
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100 velahydvilsypanthpwyisiinedgneifntsifeppppgyenvadivppfsafspq 167
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Pred. No. 0;
0; Mismatches 0;
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Synthetic.
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Inducing immune responses to weakly immunogenic, peptide antigens for the treatment of breast and
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vaccitation; cytotoxic Tymphocyte immunity; breast cancer; 
prostate cancer; cell-associated poptide antigen; foreign epitope.
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21.41
/label= P30
/note= "foreign epitope"
674.688
/label= P2
/note= "foreign epitope"
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98US-0105011.
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/note= "foreign epitope
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Prepride "Asbel- P2" Peptide "Asbel- P2" Prepride "	Example 1: Page -: 220pp: English.  MExample 1: Page -: 220pp: English.  MExample 2: Page -: 220pp: English.  MEXAMPLE 3: Page -: 220pp: English.  MEXAMPLE 4: Page -: 220pp: English.  MEXAMPLE 5: Page -: 220pp: English.  MEXAMPLE 6: Page -: 220pp: English.  MEXAMPLE 6: Page -: 220pp: English.  MEXAMPLE 7:	7
Query Match  95.0%; Score 638; DB 21; Length 750;  Best Local Similarity 100.0%; Piede 100; O; Mismatches  Qy 1: SSELTHIPPHINIAGH 100; O; Mismatches  Qy 1	De 26 everylphiphyphyphyphyphyphyphyphyphyphyphyphyphyp	Mon Apr 9 11  QY 181  Db 225

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                                                                                                                                    Homo sapiens.
Synthetic.
                                                                                                                                                        Prostate specific membrane antigen; immunogenized construct; mutant; vaccination; cytotoxic T-lymphocyte immunity; breast cancer; prostate cancer; cell-associated peptide antigen; foreign epitope.
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antigen (PBM) constructs, which contain foreign epitopes (PS and/or P30)
The immunogenic analogues of PBM can be used in the claimed method as an
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Gautam A, Birk P, Karlsson G;
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Synthetic
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                                                                                       Location/Qualifiers
21.41
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606.620
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24.38
/label= P2
/note= "foreign epitope"
607.627
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Matches 561; Conservative 0;
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Gautam A, Birk P, Karlsson G;
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Haaning J, Leach D, Dalum I;

Inducing immune responses to weakly immunogenic, peptide antigens for the treatment of breast and tumor associated prostate cancer

AX 192627-49 are mutant immunogenized human prostate specific membrane of tigen (PSM) constructs, which contain forzign epitopes (PZ and/or PSO).

The tigen (PSM) constructs, which contain forzign epitopes (PZ and/or PSO).

The tigen (PSM) contained the prostate property of the contained method as an activament of the prostate property.

The prostate property of the prostate involved in the claimed method as an activate property of the continuous pr

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99627-9 are mutant immunogenized human prostate specific meshrane antisen (PSM) constructs, which contain forting epitopes (PS and/or PSO). The immunogenic analogues of PSM can be used in the chained method as an autovaccine to induce a CTI response. Subdominant CTI pitopes, antibody binding regions and cysteine residues involved in disultide bonds are preserved in the immunogenized forms. The method is used for indusing immune responses against weakly immunogenic cell associated pertide antigens (PS) such as those associated with concers (self-proteins). e.g. unann prostate specific membrane suntigen (PSM), heregina PSM proteins (PSM) and pSM) a
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20-0CT-1998;
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                                                                                                                                                                                                                                                                                                                         Inducing immune responses to weakly immunogenic, peptide antigens for the treatment of breast and
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Synthetic.
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21.41
/label= P30
/note= "foreign epitope"
213.27
/label= P2
/note= "foreign epitope"
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98US-0105011.
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/note= "foreign epitope"
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group derived from the PA and/or at least 18 cell group derived from the cell-associated PA and (2) at least 1 first 1 helper cell group which is dreshing to the animal. Analogues of human PSM, human Herz and human/murine PSPB comprising a substantial part of all known and predicted CTL and 8 cell epitopes of the respective PA and including at least one foreign 7 helper epitope are also claimed. The method is used to treat prostate, prostate/breast or breast cancer when the PA is human RSM, RPSB and Herz), respectively.
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Sequence 750 AA;

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Query Match 74.2%; So
Best Local Similarity 100.0%; P
Matches 524; Conservative 0;
Score 524; DB 21; Length 750;
Pred. No. 0;
0; Mismatches 0; Indels 0;
Gaps
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 243
 VGLDSIPVBPTCYTAAGKLEBMOGOSLPOSSWIGGLKYPKYMOGOFYCARSTOCKWALI 302

 287
 vglpsipvhptgydadkllekmggacapdasavrgsikvpynvgpgftgntstckkkmhi 346
 183 LYSDPADYENDEVRSYPPOCHNIEGGGYORGHLINIAGGDFJRGYPANETAYRRGIABA 242 227 Lysdpadyfapgrksypddynnlygggrqrgnilnlingsddplicpyypnnyyyrsglaca 286

483 RLGIJASGRANYTKHNETHKESCYPLYSVYETYELVEKYNDPKFKHLTVAQVRGANYEE 527 rlgiasgrarythnwethkisgyplyhavyetyelvekiydpmikyhltvaqvrggmvie 586

54 B 54 B 53

663 GIYDALFDIESKYDPSKAWGEVKRQIYVAAFTVQAAAETLSEVA 706

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co antigen (PSM) constructs, which contain foreign epitopes (PS and/or PD).

Contraction to induce a CPL response. Subdominant CPL epitopes, antibody continuing a property and property and the contraction of the contractio
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64 B 

488 SGRARYTKNWETNKFSGYPLYHSVYETYELVEKFYDPWFKYHLTVAQVRGGNVFELANSI

Sequence 750 AA;

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Gaps

Ouery Match 73.5%; Score 519; DB 21; Length 750; Best Local Similarity 100.0%; Pred. No. 0; Matches 519; Conservative 0; Mismatches 0; Indels 0; Indels 0;

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22

Y92649 standard; Protein;

693

10-AUG-2000 (first entry)

Mutant human PSM antigen splice variant construct, hPSM'10.3.

Prostate specific membrane antigen; immunogenized construct; mutant; vaccination; cytotoxic T.lymphocyte immunity; breast cancer; PSM; prostate cancer; cell-associated peptide antigen; foreign epitope.

Homo sapiens Synthetic

Key Peptide

Peptide Location/Qualifiers
153..173
/label= P30
/note= "foreign epitope"
517..6abel= P2
/note= "foreign epitope"

WO200020027-A2.

05-OCT-1999; 13-APR-2000. 98DK-0001261 98US-0105011 99WO-DK00525

05-0CT-1998; 20-0CT-1998; (MEBI-) M & E BIOTECH AS.

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Page 69

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Page 70

Y92635 standard; Protein; 750

RESULT 21 Y92635 ID Y92635 XX AC Y92635 XX

Mutant human prostate specific membrane antigen construct, hPSM1.3. 10-AUG-2000 (first entry)

Prostate specific membrane antigen; immunogenized construct; mutant; vaccination; cytotoxic T-lymphocyte immunity; breast concer; prostate cancer; cell-associated peptide antigen; foreign epitope.

Homo sapiens. Synthetic.

Key Peptide Peptide Location/Qualifiers
24.38
/label= P2
/note= "foreign epitope"
210..230
/label= P30
/note= "foreign epitope"

WO200020027-A2

13-APR-2000.

98DK-0001261 99WO-DK00525

05-OCT-1998; 20-OCT-1998; 05-OCT-1999; (MEBI-) M & E BIOTECH AS.

Steinaa L, Mouritsen S, Nielsen KG, Haaning J, Leach D, Dalum I; Gautam A, Birk P, Karlsson G;

WPI; 2000-349917/30.

Inducing immune responses to weakly immunogenic, tumor associated peptide antigens for the treatment of breast and prostate cancer

Example 1; Page -; 220pp; English.

Y92627-49 are mutant immunogenized human prostate specific membrane

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Steinaa L, Mouritsen S, Nielsen KG, Gautam A, Birk P, Karlsson G;
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                                                                                                               J, Leach D, Dalum
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Page 74

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**4**26

Example 1; Page ~; 220pp; English. Inducing immune responses to weakly immunogen peptide antigens for the treatment of breast i WPI; 2000-349917/30. and tumor associated prostate cancer

A 9927-49 are mutant immunogenized human prostate specific membrane and program of the membrane of the program of the program

Sequence 693 AA;

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                                                                    Query Match 62.7%; Sometime 100.0%; I Best Local Similarity 100.0%; I Matches 443; Conservative 0;
Score 443; DB 21; Length 693;
Pred. No. 0;
0; Mismatches 0; Indels
                                                                    Indels 0;
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A Y9327-49 are mitant immunogenized human prosate specific ambotane antigen (SSM) constructs, which contain Cartain epitopes (PE and/or PSO). The immunogenite analogues of PSN con he used in the chimed mathod as not controlled to the chimed the state of the chimed action as an extraorchize to induce a CTL response. Subdeminant CTL epitopes, antibody CD binding regions and cysteine residues involved his district periods antibody CD binding regions and cysteine residues involved his cancer cell reproteins), e.g. Commune responses against weakly immunogenic cell-associated peptide contigens (PSN), and as those associated with cancers (eslf proteins), e.g. Chuman protate specific membrane antigen (PSN), hereguin 2 (Her2) and/or CD bibroblast growth factor Bb (FSRb). The method comprises effecting calmalantament system of: (1) at least 1 CTL (cyotoxia T-lymphocyte) of the calmalantament system of: (1) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyotoxia T-lymphocyte) of containants and (2) at least 1 CTL (cyot
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Gautam A, Birk P, Karlsson G;
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20-OCT-1998;
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                                                                         Homo sapiens.
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Peptide
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                                                                                             Prostate specific membrane antigen; immunogenized construct; mutant; vaccination; cytotoxic T-lymphocyte immunity; breat cancer; prostate cancer; cell-associated peptide antigen; foreign epitope.
                                                                                                                    Mutant human prostate specific membrane antigen construct, hPSM10.3
                                                                                                                                  10-AUG-2000
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210..200
/label= P30
/note="foreign epitope"
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591 IVLPFDCRDYAVVLKYAKKIYSISMKHQDBMTYSVSPDSLESAVKHPELKAKSEBL

591 IVLPFDCRDYAVVLKYAKKIYSISMKHQDBMKTYSVSfdSlfsavknfteiaskfseil
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531 ASGIALYKNWETKKEGYPLHSVYETYELVEKYYDHEKYHLIYNQVSGGWYELANS
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        aipwhpigyydaqkilekmggaappdsswrgalkvpynvgpgftgnfarqkvkmhlhatn

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Location/Qualifiers
21..41
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305..319
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Prostate specific membrane antigen; immunogenized construct, mutant; vacchantion; optotoxic T-lymphocyte immunity; breast cancer; prostate cancer; cell associated peptide antigen; foreign epitope.

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Query Match 62.7%; Sco Best Local Similarity 100.0%; P Matches 443; Conservative 0;

Score 443: DB 21: Length 750; Pred. No. 0: 0: Mismatches 0: Indels 0:

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                   A v92627-49 are nutant immunogenized human prostate specific membrane considering (FBM) constructs, which contain foreign epitopes (FP and/or F30). The very serious considering the consideri
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20-0CT-1998;
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Peptide
Y99637-49 are matant immunogenized human prostate specific membrane antiesn (PSM) constructs, which contain Carting spikopes (PS and/or PSM). The immunogenize antalogues of PSM can be used in the claimed method as an autoveccine to induce a CTI response. Subdominant CTI spitopes, antibody binding regions and cysteine residues involved in disulfide bonds are preserved in the immunogenized forms. The method is used for inducing immune responses against veakly immunogenize cell-associated peptide antigens (PS) such as those associated with concers (self-proteins). e.g. thuman protate specific membrane antigen (PSM), heregulin 2 (Herz) and/or throblast growth factor obl (PSPM). The method comprises effecting
                                                                                                                                                                                                                                               Steinaa L, Mouritsen S, Nielsen Gautam A, Birk P, Karlsson G;
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Synthetic.
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Gautam A, Birk P, Karlsson G;
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                                                                                                                                                                             Inducing immune responses to weakly immunogenic, tumor associated peptids antigens for the treatment of breast and prostate cancer
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24..38
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Query Match 57.1%; Sometive 100.0%; I Matches 403; Conservative 0;
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Matches 398: Conservative
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121 SPOGNPECDLYPVBYARTEDFFKLERDMKINGSKTVIARYGKNFRGKHVBNOLAGAG
165 SPOGNPEGDLYPVBYARTEDFFKLERDMKINGSBKIVIARYSKVFTSDKVKNAGLAGAKG
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192627-49 are mutant immunogenized human prostate specific membrane antigen (PSH) constructs, which contain foreign epitopes (P2 and/or P30). The immunogenic analogues of PSH can be used in the claimed method as an autovaccine to induce a CTL response. Subdominant CTL epitopes, antibody
                                                                                                                            Example 1; Page -; 220pp; English.
                                                                                                                                                                               Inducing immune responses to weakly peptide antigens for the treatment.
                                                                                                                                                                                                                                                                                                            Steinaa L, Mouritsen S, Nielsen
Gautam A, Birk P, Karlsson G;
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Peptide
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Key
Peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-0CT-1998;
20-0CT-1998;
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vaccination; cytotoxic [7]apphocyte immunity; breast cancer; PSM;
prostate cancer; cell-associated peptide antigen; foreign epitope.
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606.60
/label= P2
/label= foreign epitope*
210.230
/label= P30
/note= foreign epitope*
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153..173
/label= P30
/note= "foreign epitope"
549..563
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487 ASGARYTKHETHKESGYPLHENYETYELVERKYDDHEKHLTYAQVRGGMYFELANS
474 ABGRERYLKHWEINKESGYPLYBBNYG-YD-LVEKYDDHEKYDLTVAGYTSGYMCELANS
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Peptide
                                                                                                                                                                                                                                                    Steinaa L, Mouritsen S, Nielsen KG, Gautam A, Birk P, Karlsson G;
                                                                                                                                                                                                                                                                                                                                             05-OCT-1998;
20-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Key
Peptide
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Gautam A, Birk P, Karlsson G;
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20-OCT-1998;
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Synthetic.
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vaccination; vytotoxic T-lymphocyte immunity; beast cancer; PSK;
prostate cancer; cell-associated peptide antigen; foreign epitope.
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972627-49 are mutant immunogenized human prostate specific membrane antigen (PSM) constructs, which contain foreign epitopes (P2 and/or P10). The immunogenic analogues of PSM can be used in the claimed method as an attowaccine to induce a CTL response. Subdominant CTL epitopes, antibody binding regions and cysteine residues involved in disulfide bonds are preserved in the immunogenized forms. The method is used for inducing
                                                                                                                                                                                                                                                                                                                                                                                                  05-OCT-1999;
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Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Prostate apecific membrane antigen; immunogenized construct; mutant; vaccination; cytotoxic Tymphocyte immunity, breast cancer; prostate cancer; cell-associated peptide antigen; foreign epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mutant human prostate specific membrane antigen construct, hPSM6.3.
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                                                                                                                             Example 1; Page -; 220pp; English
                                                                                                                                                                 Inducing immune responses to weakly immunogenic, tumor associated peptide antigens for the treatment of breast and prostate cancer
                                                                                                                                                                                                                      WPI; 2000-349917/30.
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153.173
/label= P30
/note= "foreign epitope"
391.405
/label= P2
/note= "foreign epitope"
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98US-0105011
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448..462
/label= P2
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210..230
/nate= P30
/note= "foreign epitope"
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CC The immunogenic analogues of FSM can be used in the claimed method as an automatic property of the immunogenic analogues of FSM can be used in the claimed method as an automatic property of the immunogenic property of the property of the immunogenic property of the property of the property of the property of the immunogenic property of the i
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Controllar growth factor by (PSPM). The method comprises offerching
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Controllar sociated PA; and C) at least 1 for call group derived from the
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Correlyn to the animal. Analogues of human PSM, human Herz and
Controllar for PSP comprising a substantial part of all known and
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Best Local Similarity 100.0%; I
Matches 289; Conservative 0;
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                                                    Y92624 standard;
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465 eveffqrigianguarythnwetnkfagyplyhnsvyetyelvekfydpmfkyhltvaqvrg 524
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582 gmvfelansivljefdordyavvlryadkiyəismkhpqemktysvsidsifeavknite 641
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645 gesfeyiydalfdleskvdpskawgevkrqiyvaaftvqaanetleeva 693
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0; Mismatches
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Pred. No. le-290;
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Murine prostate specific membrane antigen

splice variant.

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Length 694;

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Query Match 5.1%; Score 36; DB 21; 1 Best Local Similarity 100.0%; Pred. No. 1.4e-28; Matches 36; Conservative 0; Mismatches 0;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 5.1%; Score 36; DB 21; Best Local Similarity 100.%; Pred, No. 1.4e-28; Matches 36; Conservative 0; Mismatches 0;
                                                                   Prostate specific membrane antigen; splice variant; mutant; vaccination, cytotoxic "lymphocyte immunity; breast cancer; prostate cancer; cell-associated peptide antigen; foreign epitope
Mus musculus.
Synthetic.
                                                                                                                                                             Mutant murine PSM splice variant construct, mPSM'Y.
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N-PSDB; A09459
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Gautam A, Birk P, Karlsson G;
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20-OCT-1998;
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cytotoxic T-lymphocyte immunity; self-protein; cancer; breast cancer;
prostate cancer; cell-associated peptide antigen; foreign epitope.
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98US-0105011.
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               RESULT 33
Y92661
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                                                A y9363-62 are nutant immunogenized murine prostate specific membrane artigan. 286 miles membrane membrane miles by immunisation addorn DBA vaccination against murine pBM customers miles by immunisation addorn DBA vaccination against marine pBM customers maked an autoreaction to induce a CIT response. The membrane against weakly immunogenic cell-associated with the ceancers (self-proteins), e.g., human PBM, heregulin 2 (Herz) and/or CC fibroblast growth factor Bb (PGPB). The method comprises effecting calmalianeous presentation by antigen producing cells (ARcs) and/or cell-associated this miles are miles of: (1) at least 1 CIT (cytotoxic Tlymphocyte) group derived from the PA and/or at least 1 Br-cell group derived from the CC calman/murine FGPB comprising a substantial part of all known and comprising a substantial part of all known and comprising as substantial part of all known and comprising as substantial part of all known and comprising a substantial part of all known and c
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20-0CT-1998;
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Peptide
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               Sequence
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Gautam A, Birk P, Karlsson G;
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Synthetic
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prostate specific membrane antigen (PSM), hersgulin 2 (Her2) and/or fibroblast sporth factor Bb (PSFB). The method comprises effecting simultaneous presentation by antigen producing cells (APGs) of the animals immune system of: (1) at least 1 CFL (Cytotoxic T-lymphocyte) group derived from the PA and/or at least 1 B-cell group derived from the CO: (1) at least 1 B-cell group derived from the cell-associated PA; and (2) at least 1 first T helper cell group which is Coreign to the animal. Analogues of human PSM, human Her2 and human/marine RFBB comprising a substantial pSM, human Her2 and breakted CFL and B-cell epicopes of the respective Paned Enclaving at least the presence of the prospective and the century breakted CFL and B-cell epicopes are take ollhed PPs method is used by the prospective PSM. FGFB and Her2, respectively.

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Matches 36
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                                                                                                                                                                                                                              Murine prostate specific membrane antigen
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al Similarity 100.0%; Pred. No. 1.4e-28;
36; Conservative 0; Mismatches 0;
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Query Match Best Local S Matches 36

oh 5.1%; Sc | Similarity 100.0%; P | 36; Conservative 0;

Score 36; DB 21; Pred. No. 1.5e-28; 0; Mismatches 0;

Length

0,

Gaps 0

sequence

752 AA;

WPI; 2000-349917/30. N-PSDB; A09458. Steinaa L, Mouritsen S, Nielsen Gautam A, Birk P, Karlsson G; Pipe claims detail a method for inducing immune responses against weakly immunogenic cell-associated peptide antigens (PA) such as those associated with cancers (i.e. self-proteins), for example, human Inducing immune responses to weakly immunogenic, peptide antigens for the treatment of breast and Example 1; Page 204-206; 220pp; English. ĸ, Heaning Ģ tumor associated prostate cancer Leach Ó Dalum

PRESULT 36
Y22659
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Y22659:
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Y22659:
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Y22659:
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HNIANT MUTINE PROS
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HNIANT MUTINE PROS
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PROSTATE CANCELLAS.
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PROSTATE
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PRI

Prostate specific membrane antigen; immunogenized construct; mutant; vaccination; cytotoxic T-lymphocyte immunity; breast cancer; prostate cancer; cell-associated peptide antigen; foreign epitope.

Mutant murine prostate specific membrane antigen construct, mPSMX

(first entry)

Protein: 756

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Mon
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                                                                                                             Steinaa L, Mouritsen S, Nielsen KG, Gautam A, Birk P, Karlsson G;
                                                                                                                        (MEBI-) M & E BIOTECH AS.
                                                                                                                                                   9 11:14:10
                                                                                                                                                   2001
                                                                                                                               98DK-0001261.
98US-0105011.
                                                                                                                                                   5
                                                                                                                                                   .08-403-803-2_copy_45_750.olig
                                                                                                                 Haaning J,
                                                                                                                 Leach D,
                                                                                                                 Dalum
                                                                                                                                                   Pag
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Inducing immune responses to weakly immunogenic, peptide antigens for the treatment of breast and Example 1; Page -; 220pp; English. tumor associated prostate cancer

A 192659-62 are mutant immunogenized murine prostate specific membrane CC satisfer (PSM) constructs, which contrain a foreign splitope. PSM the cast of the cast o

Query Match 5.1%; So Best Local Similarity 100.0%; P Matches 36; Conservative 0; Score 36; DB: ; Pred. No. 1.50 0; Mismatches DB 21; I . 1.5e-28; ches 0; Length 756; Indels 0 Gaps

Sequence

756 AA:

B 5

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99WO-DK00525

Location/Qualifiers 255..275 /label= P30

Page 99

us-08-403-803-2\_copy\_45\_750.olig

37

Y92660 standard; Protein; 761 ζ

Y92660; 10-AUG-2000

Mutant murine prostate specific membrane antigen construct, mPSMY

(first entry)

Prostate apecific membrane antigen: immunogenized construct; mutant; vaccination; cytotoxic "rlymphocyte immunity; breat cancer; prostate cancer; call-associated peptide antigen; foreign epitope.

Mus musculus. Synthetic.

Key Peptide Location/Qualifiers 689..709 /label= P30

13-APR-2000. WO200020027-A2

05-OCT-1999; 99WO-DK00525.

05-OCT-1998; 20-OCT-1998; 98DK-0001261 98US-0105011

(MEBI-) M & E BIOTECH AS.

Steinaa L, Mouritsen S, Nielsen Gautam A, Birk P, Karlsson G; Š, Haaning ŗ Leach á Dalum

WPI: 2000-349917/30.

Inducing immune responses to weakly immunogenic, peptide antigens for the treatment of breast and tumor associated prostate cancer

Example 1; Page -; 220pp; English.

Y93659-62 are mutant immunogenized murine prostate specific membrane antigen (PSM) constructs, which contain a foreign epitoge, P90. The banking makes by immunisation and/or DBA vaccination against murine PSM can be used to gitted whether autoroclarance to sevue PSM can be broken as the by immunisation and/or DBA vaccination against murine PSM thing murine PSM and appropriate antigeness makes a CTL repetion be used in the contained method and contained method is used for inducing immune responses sequent weakly immunogenic cell associated peptide antigens (Ps) auch as those associated with cancers (self-proteins), e.g. human PSM, heregulin I (Mex2) and/or cancers (self-proteins), e.g. human PSM, heregulin I (Mex2) and/or

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Page 102

Modified-site

/note" "N-glycosylated"
185
/note" "N-glycosylated"
314
/note" "N-glycosylated"
449
/note" "N-glycosylated"
600
/note" "N-glycosylated"
628
/note" "N-glycosylated"

Modified-site Modified-site Modified-site Modified-site Mon

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ithroblast growth factor 8b (RGPBb). The method comprises effecting simultaneous recentration by antigent lower (Cytockor Llymphocyte) group derived from the Pa and/or at least 18 (Cytockor Llymphocyte) group derived from the Pa and/or at least 18 (Cytockor Llymphocyte) to the specific state of the particular pall group derived from the foreign to the animal. Analogues of thuman PSA humar or 19 group which is human/mutrine PSRB comprising a substantial part of all known of predicted CFI and B-cell peptopes of the respective PA and including at least one foreign T believe springe are also claimed. The method is used to treat prostate, protein-protein the respective PA and including at the proper state, protein-protein the PSRB and Herz, respectively.

Note: This sequence was constructed from the violetype murrine PSM (Y92623), which appears on pages 204-206 of the specification.
Sequence
197
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**និងខ្លួនខ្លួនខ្លួនខ្លួន** និង

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Query Ma
Best Loc
Matches
395 SRLLQERGVAYINADSSIEGNYTLRVDCTDLMYSLV 430
                                             Match 5.1%; Score 36; DB Local Similarity 100.0%; Pred. No. 1.1 Local Similarity 0; Mismatches 36; Conservative 0; Mismatches
                                               DB 21; I
. 1.5e-28;
.hes 0;
                                                                         Length 761
                                                 Indels
                                                 0
                                               Gaps
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Y58878;
         Y58878 standard;
                   38
         Protein;
         740
         ₹
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WPI; 2000-182424/16. N-PSDB; Z58312. Pangalos M, Neefs JEFM, (JANC ) JANSSEN PHARM NV 14-JUL-1998; 14-JUL-1999; WO200004157-A2

Peeters

DCG

98GB-0015284

99WO-GB02241

Claim 14; Fig 4; 95pp; English

New human M-acetylated alpha-linked acidic dipeptidases for treating neural disorders e.g. Alähelmer's disease, schizophrenia and Parkinson's disease

B 5

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Modified-site
                                   Domain
                                                                    Homo sapiens
/note= "membrane-spanning
111
/note= "N-glycosylated"
                                    Location/Qualifiers
8..31
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MALAD-see II, W-actylated alpha-linked acidic dispeptitiase; human; chromosome Ind.4 3021; profests ences; neurodeste disease; Alzheimer's disease; achtrophrenia-nics; Parkinson's disease; Alzheimer's disease; achtrophrenia-nics; Parkinson's disease; peripheral neurophrenia-nics; Parkinson's disease; acute brain injury; mentriphe eclerosis; Partipheral neurophreetive; neuroleptic; mentripheral neurophreetive; neuroleptic; anniparkinsonian; anticonvulsant; vasotropic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human peptidase NAALAD-ase II.
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The present sequence is that predicted for human N-acetylated for librality of the basis of sport and cold disputidases [1] is predicted to the basis of sport and cold disputidases [1] is predicted to have cold to the sport and present and present protein of sport with the sport and present protein of sport with the sport and present present of sport and present present as sport and present present as sport colls, transgent corporates, polypetidas, as well as sport present present and arrangement as sport colls, transgent corporates, present as sport colls, transgent corporates, present and arrangement and considerates. The sport colls considerates are useful for treating neural disorders such as Alzheimer's disease, ac useful consistency and the sport cold of the sport of of th

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Query Match 3.4%; Score 24; DB 21; I Best Local Similarity 100.0%; Pred. No. 4e-16; Matches 24; Conservative 0; Mismatches 0;

Length 740;

Indels

0,

Gaps

0

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Š R55106 standard; 39 Peptide; 19

Prostate specific membrane antigen; PSH; prostate cancer; transmembrane glycoprotein; imaging; targeting; tumour detection, antibody detection; sequencing. Prostate-specific membrane antigen peptide fragment. 11-JAN-1995 (first entry) Homo sapiens

Key Misc-difference Location/Qualifiers /label= Trp /note= "unsure, residue present only at very levels" Mot

05-NOV-1992; 05-NOV-1993; 11-MAY-1994 WO9409820-A 9208-0973337. 93WO-US10624

WPI; 1994-167129/20. Fair WR, (SLOK ) SLOAN KETTERING INST CANCER Heston WDW, Israeli RS;

Example ; Page 44; 196pp; English Prostate-specific membrane antigen and DNA encoding useful for detecting haematogenous micro-metastatic and for identifying ligands which bind to PSM Ag

it - is tumour cells

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Sequence

740 AA;

Page

103

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The inventors attempted to sequence the PBM Ag by modified gaman degradation, explicits (\$5508-107) that gave clear dominant peaks on HPDC from the dispetted PBM Ag sample were used some of the residues were unidentified, and others were present at very low levels and identified with lower confidence. Some of these peptides encoding the PBM Ag. Alone, IN-20 was identified as a partial IBM sequence. (The full PBM Ag sequence is above in B5509) and is encoded by 065520). The PBM coding sequence is above in B5509 and is encoded by 065520). The PBM coding sequence is above in B55092 and is encoded by 065520). The PBM coding sequence is above in B55092 and is encoded or of the limitating them. The protect is useful for suppressing or for eliminating them. The protect is useful to identify or purify ligands of the Ag. It is also an attractive target for Ab-directed imaging and targeting of prostatic tumour deposits.

38888888888888888888888888888 Sequence 19 AA;

Query Match 2.7%; Surface 100.0%; I Best Local Similarity 100.0%; I Matches 19; Conservative 0; Score 19; DB 15; I Pred. No. 2.3e-12; 0; Mismatches 0; Length 19; Indels 0

Gaps 0

Š. В 40

05-NOV-1996 (first entry) W02244:

W02244 standard;

Peptide; 19

ζ

Prostate-specific membrane antigen peptide 2T46 8.

Prostate-specific membrane antigen; PSM; prostate cancer; metastasis; therapy; diagnosis.

Homo sapiens

23-FEB-1996; 29-AUG-1996 W09626272-A1

RESULAT WOODS ACC NO CONTROL OF C 02-JUN-1995; 24-FEB-1995; 02-JUN-1995; 95US-0470735. 95US-0394152. 95US-0466381. 96WO-US02424

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Best Local Similarity 100
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Prostate specific glandular kallikrin; hK2 protain; humar protate specific antigen; PBA; hK3; protate carcinoma; prostate cancer; benign prostatic hyperplasia; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tryptic peptides (M02237-45) were obtd. from the human specific membrane (PSM) antigen (see also M02234) and u design primers (785795-808) utilized in the isolation clone (785785) coding for PSM. The isolated cDMs and F are useful in developing methods for the diagnosis and of prostate cancer and metastasis.
The inventors attempted to sequence the PSM Ag by modified Edman degradation. Peptides (RS5098-107) that gave clear dominant peaks on HPLC from the digested PSM Ag sample were used. Some of the residues were unidentified, and others were present at very low levels and identified the lower confidence. Some of these peptides were used to design primers to carry out PCR to identify cDM clones encoding the PSM Ag. RS5102 was used to design primers (65277-30. A clone, IN-20 was identified as a partial PSM sequence. (The fill PSM Ag sequence is abovn in R55097 and is encoded by 065520). The PSM coding sequence is useful for suppressing or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-JUL-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Prostate specific membrane antigen; PBM; prostate cancer;
transmembrane glycoprotein; imaging; targeting; tumour detection;
antibody detection; sequencing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Prostate-specific membrane antigen peptide fragment
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                                                                                                                                                                                                                                                       Prostate specific membrane antigen and DNA encoding it is useful for detecting hematogenous micro-metastatic tumour cells and for identifying ligands which bind to PSM Ag
                                                                                                                                                                                                                                                                                                                                        WPI; 1994-167129/20.
                                                                                                                                                                                                                                                                                                                                                                           Fair WR,
                                                                                                                                                                                                                                                                                                                                                                                                                  (SLOK ) SLOAN KETTERING INST CANCER
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Misc-difference
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                                                                                                                                                                                                                   Example ; Page 46; 196pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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/note= "unidentified residue"
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                                                                                                                                                                                                                                                                                                                                                                           Israeli RS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "unidentified residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "hR2 cleavage site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INST CANCER
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; Pred. No. 2.3;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ₹
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.3e-12;
B 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
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d used to
n of a cDNA
d PSM protein
nd treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROPERTY OF STREET OF STRE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           음 경
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Apr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 2.5
Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 2.4
Best Local Similarity 100.
Matches 17: Conservative
                                                                                   02-JUN-1995;
24-FEB-1995;
02-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peptide 5 represents amino acid residues 196-213 of PMA.

Peptides 1-6 (see M4500 and M4500-08) were used as substrates to the examine amidolytic activity of prostate specific antisen (PMS) had of prostate specific glandular kalliverin (PMS) had see M45395) Peptide 6 contained 1 cleavage site for PSA and 2 for NF2 in all peptides tested, NF2 had specificity only for selected arginines and PSA primarily for selected tyrosine, phnylalanies and leucine residues. The investion provides had spootic methods using antibodies which bind prostate antigens NF2 and pro NF2, useful e 9, for monitoring the treatment or progression of prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                modulating the metastatic ability of prostate tumour cells to grow, or for eliminating them. The protein is useful to identify or purify ligands of the Ap. It is also an attractive target for Ab-directed imaging and targeting of prostatic tumour deposits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Grauer L,
Young CYF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-JAN-1998.
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        Fair WR,
                                                                                                                                                              23-FEB-1996;
                                                                                                                                                                                                 29-AUG-1996.
                                                                                                                                                                                                                                        W09626272-A1
                                                                                                                                                                                                                                                                                                 Misc-difference
                                                                                                                                                                                                                                                                                                                               Rey
Misa-difference
                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Prostate-specific membrane antigen; metastasis; therapy; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Prostate-specific membrane antigen peptide 2T34 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-NOV-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          W02241;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W02241 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 7; Fig 16; 100pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Diagnostic methods using antibodies wuseful for, e.g. monitoring treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1998-120378/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HYBR-) HYBRITECH INC. (MAYO-) MAYO FOUNDATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                W09802748-A1
                                               (SLOK ) SLOAN KETTERING INST CANCER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        43
           Heston WDW, Israeli RS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Klee GG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18 AA;
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                                                                                950S-0470735.
950S-0394152.
950S-0466381.
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                                                                                                                                                              96WO-US02424
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15..16
/note= "hK2 cleavage
                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                  /note-
                                                                                                                                                                                                                                                                                                                /note= "unidentified amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mikolajczyk SD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.5%;
70.0%;
0;
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10.0%;
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                                                                                                                                                                                                                                                                                  "unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "PSA cleavage site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 17; DB; Pred. No. 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 18; DB 19; 1
Pred. No. 2.4e-11;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ζ
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at or progression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                         PSM: prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       site"
                                                                                                                                                                                                                                                                               amino
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 15; I
3.1e-10;
ches 0;
                                                                                                                                                                                                                                                                                                                acid.
                                                                                                                                                                                                                                                                                  acid*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 18;
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Best Local Similarity 100.0%;
Matches 17; Conservative
            ¥30346;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-JUN-1995;
24-FEB-1995;
02-JUN-1995;
                                             Y30346 standard; peptide;
                                                                                                                                                   Tryptic peptides (M02237-45) were obtd. from the human prostate-
specific membrane (PSM) antigen (see also W02234) and used to
design primers (787695-809) utilised in the isolation of a cDM
clone (785785) coding for PSM. The isolated cDM and PSM protein
are useful in developing methods for the diagnosis and treatment
of prostate cancer and metastasis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 49; 284pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA encoding alternatively spliced prostate-specific membrane antigen - useful to develop prods, for detecting hematogenous micrometastic tumour cells, or prostate cancer progression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-402365/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fair WR, Heston WDW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SLOK ) SLOAN KETTERING INST CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-FEB-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Prostate specific membrane antigen; PSW; prostate cancer; transmembrane glycoprotein; imaging; targeting; tumour detection; antibody detection; sequencing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prostate-specific membrane antigen peptide fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-JAN-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tryptic peptides (80227-45) were obtd. from the human prostate-apecific membrane (PRM) antigen (see also M0224) and used design primers (736795-808) utilised in the isolation of a CNN close (736785) coding for PRM primers I-U_f1801-40). The based on amino acids 14-19 of peptide 2734 6 (M02241). The isolated CDNN and PRM process are useful in developing methods for the diagnosis and treatment of prostate cancer and metastasis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DRN encoding alternatively spliced prostate-specific membrane antigen - useful to develop prode for detecting haemacogenous micrometastic tumour cells, or prostate cancer progression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9 11:14:10 2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1996-402365/40
                                                                                                                                                                                                                                                                                                                                           15 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   950S-0470735.
950S-0394152.
950S-0466381.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Israeli RS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                    Score 15; DB 17; I
Pred. No. 2.6e-08;
0; Mismatches 0;
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Pred. No. 3.1e-10;
0; Mismatches 0;
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W02243
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AC W02243
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                                                                                Query Match 2.1%; 50
Best Local Similarity 100.0%; I
Matches 15; Conservative 0;
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Best Local Similarity 100.0%, Pred, 80. 2.6e-08
Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                 The present sequence is derived from the N-terminal of the PSM' protein, with an additional Cyg at the C-terminal. PSM' is a splice variant of pross an additional Cyg at the C-terminal. PSM' is a splice variant of pross an additional content of the Company of the of t
                                                                                                                                                                                       Sequence 16 AA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New PSM' protein, splice variant of prostate-specific membrane antigen, used as marker for prostatic cancer for diagnosis, prognosis and treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9943710-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PSM' protein; splice variant; prostate specific cancer marker; prostatic cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Prostate-specific membrane antigen; PSM; prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Prostate-specific membrane antigen peptide 2T38 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The inventors attempted to sequence the PSM Ag by modified Edman degradation. Peptides (RSSS99-10) that gave clear dominant peaks on HPG from the digested PSM Ag sample were used. Some of the residend elementation of the residend elementation of the season at very local residence of the peaks of the season at very local residence of the peaks of the season at very local residence of the peaks of the season at very local residence of the peaks 
                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 22; Page 39; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-540563/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Grauer LS, Kuus-Reichel K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BECK-) BECKMAN COULTER INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-FEB-1998;
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Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N-terminal sequence of the PSM' protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-Nov-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-NOV-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15 AA;
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1 afidplglpdrpfyr 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prostate specific membrane antigen and DNA encoding it - is useful for detecting harmatogenous micro-metastatic tumour cells and for identifying ligands which bind to PSM Ag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1994-167129/20.
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200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15
                                                                                Score 15; DB 20; I
; Pred. No. 2.7e-08;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sokoloff R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3
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                                                                                                                       Length 16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           membrane antigen
                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                             Gaps
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Page 114

Length 19; Indels

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Query Match 2.1%; Score 15; DB 20; I Best Local Similarity 100.0%; Pred. No. 3.2e-08; Matches 15; Conservative 0; Mismatches 0;

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R55103 standard; Peptide;

17

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The present sequence is derived from the N-terminal of the PSN' protein, with an additional Cys at the C-terminal and a blocked N-terminal. PSN' is a splice variant of prostute specific membrane antigen. PSN' is a marker for cancer, specifically prostatic cancer (PC). The present sequence can be used to raise monoclonal antibodies, betection of PSN' in a sample, particularly using specific antibodies, is used for disposis of prostate cancer, both in vitro and in vivo (including localization), and for prognosis of disease. Antibodies may also be compled to therapeutic spents, such as drugs, toxins, etc., and used to treat treat prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PSM' protein; cancer marker;
                                                                                                                                                                                                                                                                                                                    Claim 22: Page 39; 52pp; English
                                                                                                                                                                                                                                                                                                                                                               New PSM: protein, splice variant of prostate-specific membrane antigen, used as marker for prostatic cancer for diagnosis, prognosis and treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-540563/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Grauer LS, Kuus-Reichel K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (BECK-) BECKMAN COULTER INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-FEB-1998; 98US-0031220.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-SEP-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Y30347 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 splice variant; prostate specific membrane; prostatic cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99WO-US03810.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sokoloff R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ₹
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(SLOK ) SLOAN KETTERING INST CANCER

Israeli RS;

WPI: 1994-167129/20 Fair WR, Heston WDW,

Example ; Page 44; 196pp; English.

Prostate-specific membrane antigen and DNA encoding useful for detecting haematogenous micro-metastatic and for identifying ligands which bind to PSM Ag

tumour o cells

05-NOV-1992; 11-MAY-1994

9205-0973337 93WO-US10624

05-NOV-1993; W09409820-A Homo sapiens Prostate appeditio membrane antigen; PSM; prostate cancer; transmembrane plycoprotein; imaging; targeting; tumour detection; antibody detection; sequencing.

Prostate-specific membrane antigen peptide fragment

11-JAN-1995

(first entry)

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Αpr
Query Match 2.0%; Score 14; DB 15; Best Local Similarity 100.0%; Pred. No. 3.2e-07; Matches 14; Conservative 0; Mismatches 0;
                                                                 encoding the PSM AG. RS5103 (or RS5104) was used to design primers 065525 and 065526. A clone, IN-20 was identified as a partial PSM of Sequence: (The full PSM Ag sequence is shown in RS5097 and is encoded by 065520). The PSM coding sequence is useful for suppressing or modulating the metastatic ability of prostate timour cells to grow, or for eliminating them. The protein is useful to identify or purify ligands of the Ag. It is also an attractive target for Ab-directed imaging and targeting of prostatic tumour deposits.
                                                                                                                                                                                9 11:14:10
                                                 17 AA:
                                                                                                                                                                                                                                                                               19 AA;
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                                                                        The inventors attempted to sequence the PSN Mg by modified Edman degradation. Peptides (RS598-107) that gave clear dominant peaks on HPLC from the digested DSN Mg sample were used. Some of the residues were unidentified, and others were present at very low levels and identified with lower confidence. Some of these were used to design primers to carry out PCR to identify cDNA clones were used to design primers to carry out PCR to identify cDNA clones
us-08-403-803-2_copy_45_750.olig.ras
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 1.8%;
Best Local Similarity 100.0%;
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tryptic peptides (M922)7-45) were obtd, from the human prostate-
specific membrane (PSM) antigen (see also M922)4) and used of
design primers (758/95-808) utilised in the isabation of a CDM
clone (758/95) odding for ESM, primers E and F (756/95-800) are
based on amino acids 8-13 of peptide 2725 a (M92242). The isolated
CDMA and PSM protein are useful in developing methods for the
dispnosis and transment of prostate cancer and metastasis.
       Key
Misc-difference
                                                                                         Prostate-specific membrane antigen; metastasis; therapy; diagnosis.
                                                                                                                                       Prostate-specific membrane antigen peptide 2T47 7.
                                                                                                                                                                      05-NOV-1996 (first entry)
                                                                                                                                                                                                        W02245;
                                                              Homo sapiens.
                                                                                                                                                                                                                                     W02245 standard; Peptide;
                                                                                                                                                                                                                                                                                                                Example 1; Page 49; 284pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA encoding alternatively spliced prostate specific membrane antigen - useful to develop product for detecting heemstrogenous micrometastic tumour cells, or prostate cancer progression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fair WR, Heston WDW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1996-402365/40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            17 AA;
/note= "some uncertainty over identification
                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Israeli RS;
                                                                                                                                                                                                                                     22
                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                            Score 13: DB 17: Length 17
Pred. No. 3.4e-06;
O; Mismatches O; Indels
                                                                                                          PSM; prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                         Length 17;
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Prostate-specific membrane antigen; PSM; prostate cancer; metastasis; therapy; diagnosis. Prostate-specific membrane antigen peptide 2T35 2. 05-NOV-1996 (first entry)

W02242 standard; Peptide; 17 AA

Misc-difference Key Misc-difference

/label= Gly, Pro 15 /label= Asp, Gly

Location/Qualifiers

W09626272-A1

02-JUN-1995; 24-FEB-1995; 02-JUN-1995;

95US-0470735. 95US-0394152. 95US-0466381.

23-FEB-1996; 29-AUG-1996.

96WO-US02424

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Kisc-difference 14

Misc-difference 15

Misc-difference 15

Misc-difference 16

Misc-difference 16

Misc-difference 16

Jose Gln at position 16 is present, but at very love the servery love tha
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24-FEB-1995;
02-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Miso-difference 21

/note= "some uncertainty over identification of Glu at position 21 (sequence continues after residue 21 but at levels below the detection limit)"
                                                                       Tryptic peptides (M022)7-45) were obtd. from the human prostate-
accific measure (PMN, antique, measure who M02214) and used of the second of the control of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Misc-difference 20
/note= "Glu at position 20 is present, but at very
low levels"
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 49; 284pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DRA encoding alternatively spliced prostate-specific membrane
antigen - useful to develop prods. for detecting hamatogenous
micrometastic tumour cells, or prostate cancer progression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1996-402365/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fair WR, Heston WDW, Israeli RS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SLOK ) SLOAN KETTERING INST CANCER RES
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22 AA;
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95US-0394152.
95US-0466381.
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1.8%; Score 13; DB 17; Length 22;

Query Match

Qy 371 TILFASWDAEEFG 383 ||||||||||||| Db 1 tilfaswdaeefg 13

Best Local Similarity 100.0%; Pred. No. 4.3e-06; Matches 13; Conservative 0; Mismatches 0; Indels 0; Gaps

0;

Search completed: April 6, 2001, 16:56:36 Job time: 195 sec